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(54) Title: ANALOGS OF GLYCOPROTEIN HORMONES HAVING ALTERED IMMUNOLOGICAL CHARACTERISTICS, EFFICACY AND/OR RECEPTOR SPECIFICITY

#### (57) Abstract

The invention relates to modified chorionic gonadotropin (CG) alpha, beta-heterodimeric polypeptides comprising a gly-coprotein hormone alpha-subunit polypeptide and a non-naturally occurring beta-subunit polypeptide. The modifications in the beta-subunit polypeptide involve the deletion, addition, or substitution of one or more amino acid residues. The modified heterodimeric polypeptides of the invention have different properties compared to native hCG. Certain alpha, beta-heterodimeric polypeptides bind to LH and FSH receptors and stimulate steroidogenesis in testicular and ovarian cells. Other alpha, beta-heterodimeric polypeptides bind to LH receptors but have lower efficacy than hCG to stimulate steroidogenesis in testicular and ovarian cells.

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ANALOGS OF GLYCOPROTEIN HORMONES HAVING ALTERED IMMUNOLOGICAL CHARACTERISTICS, EFFICACY AND/OR RECEPTOR SPECIFICITY

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# FIELD OF THE INVENTION

The invention relates to glycoprotein hormone analogs and other proteins, their methods of use and other applications. The molecules of the invention are analogs or derivatives of naturally occurring vertebrate glycoprotein hormones and DNA molecules encoding them. typically occurring hormones are naturally The constituted of an alpha and beta subunit. Amino acid sequences of naturally occurring alpha and beta-subunits from several different species are presented in TABLE I. Monomeric forms of both subunits as well as heterodimers In the molecules of and homodimers are known to occur. the invention the chemical structure of the beta-subunit or the alpha-subunit or both may be different from that of the native hormones. The glycoprotein hormone analogs of the invention also differ from the naturally occurring physiological, biological, in their characteristics. immunological pharmaceutical and/or Observations that many different changes can be made in the subunits of naturally occurring glycoprotein hormones which can selectively alter specific properties of the of hormone led to increased knowledge molecules has The hormone derivatives of the invention are structure. endocrine various with which tools as useful metabolic functions can be manipulated and dysfunctions One aspect of the invention relates to can be treated. the control of reproduction in mammals.

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The hormones of the invention are useful in numerous applications including therapeutic, immunological, diagnostic, and other uses. Certain of the hormones of the invention are capable of altering the

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reproductive process in living organisms such as mammals and other vertebrates. Certain of the hormones promote fertility; others are contragestives. The various aspects of the invention are discussed in full detail hereinafter. The hormones of the invention may be produced by genetic engineering, chemical synthesis and/or a combination of such methods. The glycoprotein hormones of the invention contribute to fulfilling an important and actual need in the relevant field of art.

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### BACKGROUND OF THE INVENTION

15 The glycoprotein hormones include the gonadotropins and thyrotropins. In humans these chorionic gonadotropin (CG), luteinizing hormone (LH), follicle stimulating hormone (FSH) and thyroid stimulating hormone (TSH). They family of are a 20 heterodimeric proteins which share a common alpha-subunit differ in their hormone-specific beta-subunits. Gonadotropins such as CG, LH and FSH play a major role in the reproductive process, while TSH is important for thyroid function. Owing to the importance of these hormones in endocrine and metabolic function, there has 25 considerable interest in characterizing relationship between hormone structure and function.

While primary structures of many glycoprotein hormones have been known for many years, the relationship between hormone structure and function has been only partially characterized in spite of many studies ' involving chemical modifications (reviewed deglycosylation (1-4), immunological characterization (5-8), and use of peptide analogs (9-12). The beta-subunit specifies the activity of the hormones, however, neither free subunit has much biological activity (1). reason the alpha, beta-heterodimer is the most active form is not known. Its activity might be the result of a

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unique conformation of one or both subunits which forms after they combine (13) and/or it might be that portions of both subunits interact directly with the receptor (9, The observation that some antibodies bind to epitopes on the hCG beta-subunit only when associated with the alpha-subunit suggests that the conformation of the <u>beta-subunit</u> is altered when combines with the <u>alpha</u>-subunit (14).Residues responsible for the binding of these antibodies have been identified and include those between Cys-38 and Cys-57 (15), a region of the molecule which has been proposed to interact with the receptor (10).Similarly, monoclonal antibodies and polyclonal antisera to the alpha-subunit (16) have higher affinity for the alpha, than for the free alpha-subunit, beta-heterodimer suggesting that the conformation of the alpha-subunit is altered when it combines with the beta-subunit. Evidence that both subunits participate in binding to the receptor is based on the ability of synthetic alpha and betasubunit peptides to inhibit hCG binding to receptors (9, 10) and on the abilities of both subunits to be crosslinked to the receptor (17-19). It appears that the have a different conformation in alpha-subunit may different hormones since anti-alpha-subunit antibodies have been found which bind to the alpha-subunit in the but heterodimeric forms of some not all human glycoprotein hormones (29).

The Asn-linked carbohydrate residues influence efficacy of the hormones. Deglycosylation by chemical 30 (3, 4), enzymatic (20, 21), or genetic (22) procedures renders them less active in vitro. The reasons the carbohydrate residues are required, particularly those on the alpha-subunit at Asn-52 (22), are unknown. carbohydrate residues may participate in cross-linking 35 complex to other cell hormone-receptor proteins (2, 23). They may also alter the conformation of the molecules (24) although this has been hard to demonstrate spectroscopically (25). The carbohydrates

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may increase the flexibility of the hormones and facilitate a change in conformation which occurs after the hormones bind to the receptor (26). A report that Fab fragments of antibodies to hCG can convert the deglycosylated hormone into an agonist (27) lends further support for the effects of the carbohydrates on the conformation of hCG. Asn-linked sugars also appear to have a role in dimer stability since it is more difficult to dissociate the dimer following deglycosylation (28) and since deglycosylated subunits recombine faster than native alpha and beta-subunits (25).

are two major barriers to the understanding of the relationship between glycoprotein hormone structure and function. First, the tertiary structures of the hormones have not been determined. hormones resisted crystallization and only recently have high quality crystals for x-ray diffraction studies of a glycoprotein hormone (hCG) been obtained (71, 72). addition, there is doubt regarding the locations of the disulfide bonds in the subunits (1, 2). Due to the difficulties anticipated in obtaining the tertiary structure of the carbohydrate chain, known to influence the activity of the hormone, even crystallographic and/or NMR approaches may not guarantee the characterization of residues which interact with hormone receptors to induce a response.

The second major impediment to understanding the relationship between structure and function has been the difficulty of preparing analogs. Until 1985 (30), the only means of preparing analogs was by chemical or enzymatic means. The limited numbers of residues which could be specifically modified and the difficulty of removing particular N-linked sugar chains curtailed progress substantially.

Present day recombinant DNA technology permits the production of a much wider range of analogs. Several

glycoprotein hormones have now been expressed from DNA introduced into mammalian cells in culture, and analogs of some have also been expressed (22, 30-36).

The present invention utilizes DNA mutagenesis and gene expression to produce glycoprotein hormone analogs. These analogs have enabled identification of residues important for receptor binding and specificity, subunit interactions, and antibody binding.

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It has been discovered that a wide variety of changes can be made in these hormones. For example, chimeric subunits can be created which combine specific characteristics of each parent molecule such as receptor or antibody binding (TABLES II, III, IV, VI, XIII and FIGURE 11). Moreover, deletions of certain regions can result in molecules having decreased efficacy (TABLE VII and FIGURE 14).

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Identification of amino acid residues in the contribute to biological, that their hormones and/or immunological physiological, pharmacological, enabled characteristics (FIGURES 7-12, 14) has The invention production of a variety of useful analogs. provides for compounds useful for manipulation of gonad or thyroid function. Among these uses are the induction of fertility in male and female mammals, promotion of female infertility, contragestion and the in vitro and in vivo control of reproductive capacity in animals.

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#### BACKGROUND ART

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The glycoprotein hormones have been studied extensively. There has been considerable interest in the structure/function relationships of these molecules. The general references listed below reflect the current state of knowledge in this area.

PIERCE, J.G. and PARSONS, T.F., Ann. Rev. Biochem. 50, 465 (1981). This extensive review article summarizes the results of various chemically and enzymatically derived analogs of the naturally occurring glycoprotein hormones to discern the relationship between their structure and function. The authors conclude that much remains to be known about the interaction of the glycoprotein hormones with their receptors.

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RYAN, R.J. et al., Recent Progress Hormone Res. 43, 383 (1987). RYAN, R.J. et al., "The Glycoprotein Hormones: Recent Studies of Structure-Function Relationships", FASEB Journal 2, 2661 (1988). These articles summarize studies employing synthetic peptides to study glycoprotein hormone structure and function.

<u>KEUTMANN, H.T. et al., Proc. Nat'l. Acad. Sci.</u>

<u>USA</u> 84, 2038 (1987). This work concludes that the intercysteine 38-57 loop sequence in the <u>beta</u>-subunit of hCG and hLH is a determinant for expression of biological activity in these molecules.

- SCHNEYER, A.L. et al., Biochemistry 27, 666
  25 (1988). This work concludes that hFSH beta-subunit sequences 33-52 are important in receptor binding, and that the sequence TRDL (34-37) is of particular importance.
- 30 An example of work identifying the role of the subunits in determining binding specificity Strickland, T.W. and Puett, D., "Contribution of Subunits to the Function of Luteinizing Hormone/Human Chorionic Gonadotropin Recombinants", Endocrinology 109, 35 Hybrid recombinants prepared from bovine and (1981).porcine LH and human CG were used. It was observed that both the alpha and beta-subunits were found to contribute to the properties of the recombinant in several in vitro

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systems. In each case, the recombinant behaved most like the hormone from which the <u>beta</u>-subunit was derived.

For a general reference on reproductive endocrinology, see <u>Yen</u>, <u>S.S.C.</u> and <u>Jaffe</u>, <u>R.B.</u>, "Reproductive Endocrinology: Physiology, Pathophysiology and Clinical Management", 2nd edition, W.B. Saunders Company, Philadelphia (1986), hereinafter referred to as "Yen and Jaffe". Chapter 19 of this text reviews both the mechanisms involved in fertilization and the clinical problems of infertility. This text illustrates uses of glycoprotein hormones for induction of fertility.

### BRIEF DESCRIPTION OF THE FIGURES AND TABLES

FIGURE 1 depicts the nucleotide sequence and encoded amino acid sequence of the hCG <u>beta</u>-subunit cDNA along with several restriction endonuclease recognition sites.

FIGURE 2 depicts the nucleotide sequence and encoded amino acid sequence of the hCG <u>alpha</u>-subunit cDNA along with several restriction endonuclease recognition sites.

FIGURE 3 depicts the sequences DNA oligonucleotides used to construct plasmids pKBMt and pKBM.

FIGURE 4A depicts the nucleotide sequence and encoded amino acid sequence of the analog hCG-beta, which contains two silent mutations that eliminate restriction endonuclease recognition sites present in the hCG beta-DNA depicts the subunit CDNA; and FIGURE4B oligonucleotides used to construct this analog by cassette mutagenesis.

FIGURE 5 depicts the nucleotide sequences of DNA oligonucleotide cassettes used to construct analogs F8 (see TABLE VII), B9 and B11 (see TABLE II).

- FIGURE 6 depicts the nucleotide sequence and encoded amino acid sequence of the synthetic bovine <a href="mailto:alpha-subunit gene">alpha-subunit gene</a>, along with several restriction endonuclease recognition sites.
- 10 FIGURE 7 illustrates human <u>alpha-subunit</u> residues critical for antibody binding.

FIGURE 8 illustrates hCG <u>beta</u>-subunit residues critical for antibody binding.

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FIGURE 9 is a dose response curve of LH receptor binding by hCG/hFSH chimeric <u>beta</u>-subunits co-expressed in COS7 cells with human <u>alpha</u>-subunit, compared with hCG and hFSH.

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FIGURE 10 is a dose response curve of FSH receptor binding by hCG/hFSH chimeric <u>beta</u>-subunits co-expressed in COS7 cells with human <u>alpha</u>-subunit, compared with hCG and hFSH.

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FIGURE 11 is a dose response curve of LH receptor mediated steroidogenic activity of hCG/hFSH chimeric <u>beta</u>-subunits co-expressed in COS7 cells with human <u>alpha</u>-subunit, compared with hCG and hFSH.

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FIGURE 12 is a dose response curve of FSH receptor mediated steroidogenic activity of hCG/hFSH chimeric beta-subunits co-expressed in COS7 cells with human alpha-subunit, compared with hCG and hFSH.

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FIGURE 13 is a dose response curve of LH receptor binding by an hCG <u>beta</u>-subunit deletion analog (F8) co-expressed in C127 cells with human <u>alpha</u>-subunit, compared with hCG.

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FIGURE 14 is a dose response curve of LH receptor mediated adenylate cyclase activity of an hCG beta-subunit deletion analog (F8) co-expressed in C127 cells with human alpha-subunit, compared with hCG.

TABLE I depicts amino acid sequences of several naturally occurring glycoprotein hormone <u>alpha</u>-subunits and beta-subunits.

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TABLE II depicts amino acid sequences of several hCG/hFSH beta-subunit chimeras.

TABLE III depicts amino acid sequences of several hCG/hTSH beta-subunit chimeras.

TABLE IV depicts amino acid sequences of several hCG/bLH beta-subunit chimeras.

TABLE V depicts amino acid sequences of several hCG/eLH <u>beta</u>-subunit chimeras.

TABLE VI depicts amino acid sequences of several hCG/hLH beta-subunit chimeras.

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TABLE VII depicts amino acid sequences of several <a href="mailto:beta">beta</a>-subunit deletion mutants.

TABLE VIII depicts amino acid sequences of several other <u>beta</u>-subunit analogs.

TABLE IX depicts amino acid sequences of several <a href="mailto:beta-subunit-analogs">beta-subunit-analogs</a>.

35 TABLE X depicts examples of several <u>beta-</u> subunit/fusion proteins.

TABLE XI illustrates differences in monoclonal antibody binding among chimeric human/bovine <a href="https://beta-recommons.org/beta-recommons.org/">beta-recommons.org/</a>

subunits co-expressed with hCG <u>beta</u>-subunit compared with human and bovine <u>alpha</u>-subunits co-expressed with hCG <u>beta</u>-subunit.

TABLE XII illustrates the ability of membrane bound hCG-<u>beta</u>/VSV-G fusion protein to combine with the hCG <u>alpha</u>-subunit and be recognized by monoclonal antibodies to <u>beta</u>-subunit (B105), <u>alpha</u>-subunit (A111) and the <u>alpha</u>, <u>beta</u>-heterodimer (B109).

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TABLE XIII illustrates monoclonal antibody binding to chimeric hFSH/hCG beta-subunits.

TABLE XIV illustrates LH receptor binding by alpha, beta-heterodimers containing chimeric alpha-subunits or alpha-subunit deletions.

TABLE XV illustrates monoclonal antibody binding to hCG-<u>beta</u>/VSV-G fusion proteins on stably transfected COS7 cells compared with COS7 cells stably transfected with hCG <u>beta</u>-subunit.

#### SUMMARY OF THE INVENTION

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The invention provides protein hormone analogs which are "engineered" such that they are capable of controlling various hormone-regulated physiological functions in vertebrates of both genders. In particular, the invention relates to the glycoprotein hormones and their individual subunits. The heterodimeric forms of the hormones are thought to be the most active. The four human hormones belonging to this group are chorionic gonadotropin (CG), luteinizing hormone (LH), follicle stimulating hormone (FSH) and thyroid stimulating hormone (TSH).

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accordance with the invention, In terminology "hormone analogs", "glycoprotein hormone analogs" or "variants" is to be understood in the Within the description the broadest manner. "chimera" is used to designate a hormone analog which contains amino acid sequences derived from two or more These chimeras can be different glycoprotein hormones. constructed from proteins with low sequence homology. For example, beta-subunit chimeras can be made from hCG and hFSH beta-subunits in regions of very low sequence homology (TABLE II). These chimeras, described in detail below, exhibit useful properties such as the ability to bind and stimulate both LH and FSH receptors. In some instances comparatively small changes, such modification of a single amino acid in the hormone sequence, cause significant alteration of the activity as compared to the native hormone.

analogs contemplated by the invention molecules which have novel biological, 20 include physiological, pharmacological and/or immunological (hereinafter "biological") activities and molecules that intermediates for such active molecules. useful Within the scope of the invention are also modified alpha and/or beta-subunits, not only the coupled dimers. These 25 subunit analogs may be used in the monomeric ("free subunit") form. The modified subunits may also be coupled with an appropriate other subunit. Further, as will become apparent from the description, the hormone analogs of the invention need not be constituted from the 30 same animal species, they can be prepared using portions obtained combinations of the hormones from vertebrate, i.e., "cross-species" chimeras (for examples see TABLES IV, V, and IX). Further, within the scope of the invention are applications and uses in the human 35 field, in the veterinary field, and other fields of use, including applications in other vertebrates. As further described, some analogs of the invention have the

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property of influencing fertility (e.g., promoting or inhibiting) in humans and animals.

In accordance with the invention, particular domains of amino acids are identified to which specific physiological functions or other biological activities can be attributed (FIGURES 7-12). The invention encompasses a wide variety of variants of the alpha and beta-subunit which in the dimer or monomer form exhibit new or altered biological activities. It is contemplated that some analogs of the invention will be useful as agonists or antagonists which can act as receptors for one or more different glycoprotein hormones.

In another aspect of the invention, fusion proteins have been made between a glycoprotein hormone beta-subunit (hCG-beta) and another protein (TABLE X). Particularly there is described an hCG beta-subunit/VSV G-protein analog, in which amino acid residues from the transmembrane and cytoplasmic domains of the vesicular stomatitis virus G-protein have been inserted in the hCG-beta-subunit downstream of Cys-110 at hCG-beta using the SmaI restriction endonuclease site at residues 135-137 (TABLE XII). It is contemplated that this analog will be useful as a screening system for analysis of mutagenesis products and as an immunogen which can be introduced into an animal or human attached to the cells or organisms (such as a virus) producing it, thereby eliminating the need to isolate and purify the protein itself.

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The invention provides for proteins in which amino acid residues from a glycoprotein hormone subunit (or subunits) are inserted into another protein. is contemplated that it hormone residues involved in antibody recognition may be transferred to a non-glycoprotein hormone (<u>e.q.</u>, lysozyme) resulting protein used as an immunogen to elicit an response against selected regions glycoprotein hormone. Another aspect of the invention

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provides for the design of peptides useful as immunogens which induce an immune response capable of specifically inhibiting the action of one or more glycoprotein hormones. It also provides for the design of peptides useful as receptor agonists or antagonists.

It is also contemplated that regions of a glycoprotein hormone subunit, for example hCG-<u>beta</u> 39-56 or hFSH-<u>beta</u> 33-50, could be replaced with a sequence from a non-glycoprotein hormone and used as an antigen to develop antisera or antibodies to the inserted sequence.

aspect of the invention provides betasubunit chimeras that are capable of directing hormone binding to both LH and FSH receptors. These chimeras may derived from human glycoprotein hormones hCG/hFSH or hLH/hFSH) or non-human glycoprotein hormones bovine LH/bovine FSH (e.q., or equine CG) /equineFSH). It is contemplated that these chimeras will be useful for the treatment of infertility in men and women and the promotion of fertility in male sand female animals. 7.

The invention also provides alpha-subunit Among these are chimeric <u>alpha</u>-subunits analogs. comprised of sequences from alpha-subunits from different species (TABLE IX). These analogs exhibit immunological properties. It is contemplated that these analogs will be useful immunogens. Bovine/human alphasubunit chimeras have enabled the identification of particular regions of the subunit which are involved with antibody binding (FIGURE 7), and may be used to restrict or alter the immune response to an immunogen.

35 Cross-species <u>beta</u>-subunit chimeras such as hCG/bovine LH (TABLE IV) or bovine LH/equine LH are also provided by the invention. Likewise, cross-species chimeras derived from other vertebrates are contemplated. It is contemplated that these chimeras will have

veterinary applications and also be useful for mapping regions of the hormones involved in receptor binding and biological efficacy.

Another class of chimeras provided by 5 gonadotropin/thyrotropin invention includes (TABLE III). These analogs are expected to be useful for manipulating gonad or thyroid function. They will also be useful for identifying amino acid residues responsible 10 for hormone function. This information will facilitate the production of improved hormone antagonists, agonists, agonists and analogs that agonist/antagonist activity to different receptors.

15 Some constructs provided by the invention will enable development of antibodies to specific domains on glycoprotein hormones. It is expected that these antibodies will be hormone-specific, and thus will not exhibit undesirable binding to other hormones. 20 constructs include alpha and beta-subunit chimeras and analogs in which carbohydrate groups have been introduced block antigenicity of selected regions hormones. Other analogs are contemplated in which known are placed into the T-cell epitopes hormones 25 facilitate their immunogenicity. Two utilities are anticipated. In the first, it is contemplated that these constructs will be useful as immunocontragestive These immunogens are contemplated to be useful vaccines. in this regard in that the antibodies generated will be 30 high titer and specific for particular parts of the hCG molecule. The hCG-antibody complex elicited will prevent hCG from the placenta from stimulating the corpus luteum, thereby causing termination of pregnancy at an early stage. The second application contemplates controlling 35 the reproductive capacity of animals. More specifically, the immunogenic constructs of the invention contemplated to be useful for immunocastration of steers, doas and cats and other vertebrates. Additionally, the suppression of the development of the

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reproductive system of heifers is contemplated as a way to control heat in the feed lot.

The invention also provides constructs that are glycoprotein hormone receptor antagonists. It is contemplated that these antagonists will have several applications. In particular hormone antagonists can be used to terminate ectopic pregnancy, block the action of LH in polycystic ovarian disease, suppress gonadal activity during cancer chemotherapy, or block the action of stimulatory immunoglobulins on the thyroid in Graves' disease.

In summary, the invention provides analogs of glycoprotein hormones which will have a variety of biological characteristics.

### DETAILED DESCRIPTION OF THE INVENTION

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The invention includes three major embodiments. The first includes analogs of glycoprotein hormones capable of the induction of fertility in men, women and animals of both genders. The second includes analogs which will function as immunocontragestive vaccines, immunocastration agents and suppressors of reproductive system development. Embodiment three includes antagonists capable of inhibiting hormone or receptor activity.

### INDUCTION OF FERTILITY IN WOMEN, MEN AND ANIMALS

Infertility is a significant clinical, social and human health problem. Many women who desire to become pregnant fail to develop a dominant follicle and trigger the hCG surge needed for ovulation. If an ovulation is caused by gonadotropin deficiency, such as in polycystic ovarian disease, follicular maturation and

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ovulation can be induced by administration of human menopausal gonadotropin (hMG) followed by hCG. hMG is an extract prepared from the urine of post-menopausal women containing hLH and hFSH. It is sold commercially under the tradename Perganol(TM).

In women both hLH and hFSH are necessary for ovarian steroidogenesis, follicle maturation, and ovulation. During follicular development in the ovary, LH stimulates the cells surrounding the follicle and induces androgen production. Cells inside the follicle surrounding the oocyte are responsive to FSH and convert the androgens produced by the thecal and interstitial cells into estrogens.

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constructs of the invention are contemplated to be useful for the treatment of female infertility caused by gonadotropin deficiency. molecules interact with LH and FSH receptors such that when administered infertile female, to an formation and follicular maturation should be induced. Thus, the invention discloses a protein which can replace Perganol (TM), hMG or other mixtures of hLH and hFSH in the treatment of infertility.

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Preferred embodiments of the invention chimeras in which residues near the carboxy-terminal region of hFSH are substituted for the corresponding These chimeras combine with the human region of hCG. alpha-subunit to form a dimer which has biological activity. Four specific chimeras are discussed below. The sequences of these constructs are shown in TABLE II and are denoted B11, B17, B18 and B19. All of these constructs combine with the alpha-subunit to form a heterodimer which can bind LH receptors and stimulate Leydig cell steroidogenesis (20) as well as bind FSH receptors and stimulate granulosa cell steroidogenesis (37). Antibody binding to these analogs has also been

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characterized. Results of representative assays are indicated in TABLE XIII and FIGURES 9-12.

A preferred embodiment of the invention construct B18. This construct is alternately designated beta-CF94-114, where C and F indicate the pattern of hCGbeta and hFSH-beta sequences, respectively, indicate the first and last non-identical numbers residues in each region of substitution. Construct B18 is also alternately designated DG'. In this construct, residues 88-108 of hFSH-beta replace residues 94-145 of This construct combines with the alpha-subunit hCG-beta. to form a heterodimer that can bind both LH and FSH receptors and stimulate steroidogenesis (FIGURES 11 and 12).

other embodiments of the hCG/hFSH beta-subunit chimeras are constructs B11, B17, and B19. Construct B11, alternately designated beta-CF94-117 or DG, contains residues 88-111 of hFSH-beta in place of residues 94-145 of hCG-beta. Construct B17, alternately designated beta-CFCF39-58/94-117 or ABCDG, contains residues 33-52 and 88-111 of hFSH-beta in place of residues 39-58 and 94-145 of hCG-beta, respectively. Construct B19, alternately designated beta-CFCF39-58/94-113, 114Q or ABCDG', is a variant of construct B17 which combines chimeric and non-chimeric substitutions and a deletion. The amino acid sequence is the same as B17 except that a glutamine (Q) is substituted for glutamic acid (E) at position 114, and the last three residues (115-117) are deleted.

In addition to these embodiments, it is contemplated that smaller regions of hFSH-beta such as the sequence DSDSTDCTVRGLGPSY (hFSH-beta residues 88-103), TVRGLGPSY (hFSH-beta residues 95-103) or TVRGLG (hFSH-beta residues 95-100) could be inserted into appropriate regions of the hCG (or hLH) beta-subunit and that the resultant chimeric beta-subunit would direct binding of the alpha, beta-heterodimer to FSH receptors,

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eliciting a biological response in addition to the molecule's native LH receptor binding activity and efficacy.

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5 It is also contemplated that hCG-beta or hLHbeta sequences can be substituted at the carboxy terminal of hFSH, as for construct B25, and that smaller regions of hCG-beta can be substituted into hFSH-beta, as for construct B26, and that these chimeric proteins would direct binding of the alpha, beta-heterodimer to eliciting a biological response. further contemplated that these various chimeras may exhibit differing potencies of activity at the LH and FSH receptors, leading to the selection of one over the others for certain applications because of its higher or lower ratio of FSH to LH activity.

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The finding that the constructs of invention containing hFSH-beta residues in place of hCGbeta residues 94-145 will direct binding to LH and FSH receptors and stimulate steroidogenesis was an unexpected The prior art shows that other regions of the betasubunit were thought to be important for Schneyer et al. (12) indicate that the TRDL sequence of hFSH-beta (residues 34-37) are necessary for binding to FSH receptors. Subsequent studies by Santa Coloma et. al. (76) indicate that hFSH-beta residues 33-53 bind and stimulate FSH receptors. Keutmann et al. (10) concluded that corresponding region of hCG-beta and hLH-beta, residues 38-57, were necessary for binding and stimulation of LH receptors. These authors concluded that hCG-beta residues 93-101 were necessary for binding to LH receptors. If these regions of the <u>beta</u>-subunit were important for receptor specificity, chimeric hCG/hFSH beta-subunits containing sequence alterations in these regions would exhibit altered binding specificity. This is not the case. obtained with constructs B9, B15, and B16 reflect this (FIGURES 9-12). Construct B15, alternately designated

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CFC39-58 or ABC, contains hFSH-beta residues 33-52 in place of hCG-beta residues 39-59. The receptor binding activity of the alpha, beta-B15 heterodimer is virtually indistinguishable from hCG (FIGURES 9 and 10). Construct B9, alternately designated CFC94-97 or D, contains hFSHbeta residues 88-91 in place of hCG-beta 94-97 (residues corresponding to hCG-beta 98-100 are identical in hFSHbeta and hCG-beta). The alpha, beta-B9 heterodimer binds and stimulates LH receptors, but with about 30-fold less potency than hCG (FIGURES 9 and 11). The alpha, beta-B9 heterodimer exhibits no FSH receptor binding at the amounts tested (FIGURE 10). Construct B16, alternately CFCFC39-58/94-97 or ABCD, combines designated substitutions present in B9 and B16. As for B9, the alpha, beta-B16 heterodimer is about 30-fold less active than hCG at LH receptors (FIGURES 9 and 11). It is no better than hCG in binding to FSH receptors (FIGURE 10).

It is further contemplated that the chimeric constructs of the invention would be useful for the enhancement of fertility in men and animals of male gender, i.e., to stimulate spermatogenesis. Induction of spermatogenesis requires both LH and FSH. LH stimulates testosterone production by Leydig cells of the testes. FSH and the high local concentrations of testosterone produced in response to LH stimulate the Sertoli cells and thereby induce spermatogenesis. For this reason, compounds that act on both FSH and LH receptors are Further, since it is known that hCG has a desirable. much longer half-life than FSH, it would be expected that hormone analogs containing hCG/hFSH-beta-subunit chimeras would have a longer biological half-life than hFSH. These analogs would be anticipated to be more efficacious than hCG or hFSH alone or mixtures of hCG and hFSH.

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# IMMUNOLOGICAL CONTROL OF THE REPRODUCTIVE PROCESS

It is contemplated that the glycoprotein hormone analogs of the invention will permit the

development of high titer specific antisera to hCG and thus be useful as long-term anti-fertility agents. The need for antigens capable of eliciting specific antisera to glycoprotein hormones for diagnostic immunoassays of the hormones and for the development of vaccines against hCG has long been recognized.

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is anticipated that the analogs of the invention which retain regions of tertiary structure present in hCG-beta will be considerably more potent than synthetic peptides in eliciting an immune response to specific domains of hCG and be more efficacious as an immunocontraceptive vaccine. Recent studies in which carbohydrates are added to proteins have demonstrated that it is possible to reduce their antigenicity (39). it is contemplated that the addition carbohydrate residues to some regions of the hCG-betasubunit will reduce the immunological cross-reactivity between analogs of the invention and the glycoprotein hormones (in human these are LH, FSH, and A preferred embodiment of the invention is analog G9 (TABLE VIII). In this analog hCG beta-subunit residue 78 is replaced with another amino acid and residue 79 is replaced with either a serine or a threonine residue to create a glycosylation signal for Asn-77, thereby reducing the antigenicity of the major common epitope between hLH and hCG. It is not to be excluded that analogs of the invention may be immunogenic or be of immunogenicity when conjugated appropriate carrier, such as muramyl dipeptide (MDP), derivatives thereof or other suitable carriers. The carrier may or may not be immunogenic. The immunogenic analogs of the invention are formulated with a suitable vaccine adjuvant, as is known in the art. Methods of vaccination for humans and animals are known in the art.

The antigens of the invention will generate an immune response to specific domains of the hCG molecule. The antibodies generated will bind endogenous circulating

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hCG produced by the placenta, blocking it from binding corpus luteum LH receptors with the result that the corpus luteum will stop secreting progesterone. Because of the need for ovarian progesterone in maintenance of early pregnancy, a lack of this steroid hormone will result in termination of the pregnancy.

It is also contemplated for the constructs of controlling invention useful for the to be reproductive capacity and secondary sex characteristics One such use is as an immunological animals. replacement for surgical castration of bulls. castrated to prevent them from fighting and damaging each other. However, there are numerous problems which result There is a fairly high chance of from castration. infection and some morbidity. The resultant steer also does not eat regularly for a period after surgery and as Even after an animal a result, weight gain is retarded. has recovered from surgery, the rate of weight gain remains lower and the meat produced by the animal has a higher fat content. To compensate for these effects, farmers feed their cattle androgen supplements. The in circulating androgen leads to increase faster weight gain and meat with a higher protein However, this practice has been found not to be totally acceptable. Constructs of the invention would be immunogens for generating antibodies as specific structural domains of bovine LH. The antibodies would partially neutralize the hormone resulting prevention of puberty but still provide sufficient circulating testosterone to have the positive effects on weight gain. Further, it can be seen that immunogens of suppressing be useful for invention can development of the reproductive system to control heifer in the feed lot, spay cats and dogs, preclude gelding of horses and other such applications.

Numerous other practical uses and applications of the immunogenic constructs are considered within the

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scope of this invention. The generation of high titer immune responses to particular domains of glycoprotein hormones in which the antibodies produced are not crossreactive to other glycoprotein hormone species permits the control of the physiological function with which the particular glycoprotein hormone is associated.

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#### HORMONE ANTAGONISTS

10 In accordance with the invention, constructs of the invention are useful for the non-surgical termination of pregnancy and ectopic pregnancy. Contragestion is accomplished by certain constructs of the invention which block access of native hCG to the corpus luteum. 15 constructs exhibit hCG receptor binding specificity but a decreased efficacy of steroidogenesis induction. decreased induction of progesterone formation results in inadequate levels for maintenance of the pregnancy.

A preferred specific embodiment of the hCG antagonists is construct F8 (TABLE VII). In this construct, residues 47-50 were deleted from native hCGbeta and residue 51 was changed from alanine (A) to proline (P). When combined with alpha-subunit this construct has been shown to bind LH receptors (FIGURE 13) and stimulate hCG induced cyclic-AMP accumulation at approximately 50% of the efficacy of hCG (FIGURE 14). This analog has also been found to inhibit hCG-induced cyclic-AMP accumulation (FIGURE 14). This reduces the steroidogenic potency of the analog.

It is contemplated that the hCG antagonists would be useful for suppression of gonadal activity during chemotherapy. It is further contemplated that TSH inhibitors would be useful as therapeutics in Graves' disease and that LH inhibitors would be useful for treatment of polycystic ovarian disease.

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### OTHER CHIMERAS

The invention also relates to other glycoprotein hormone chimeras or analogs. It is contemplated that these chimeras will be useful for structure/function residue mapping, immunological purposes, antagonists, contragestives, and fertility-influencing compounds for various animal species.

One important embodiment of the invention is 10 beta-subunit species alpha or analogs. cross Human/bovine alpha-subunit analogs are shown in TABLE IX. These combine with hCG beta-subunit and bind LH receptors (TABLES XI, XIV). It is contemplated that these chimeras will be useful for construction of hormone dimers that 15 can be used as an immunogen such that antibodies crossnaturally occurring glycoprotein reactive with other hormones will not be produced. It is also contemplated that cross-species chimeras may provide alternative 20 sequences to place in human hormones which may further alter the properties of the molecule.

the invention Another embodiment of subunit chimeras demonstrating cross-species hCG/bovine LH and hCG/equine LH (or CG) beta-subunit Such constructs are shown in TABLE IV and V, chimeras. It is contemplated that these constructs respectively. will be useful for identifying residues which important for binding to the human receptor. also have applications as immunogens, agonists and/or antagonists.

Another embodiment of the invention includes hCG/hTSH chimeras. Such constructs are shown in TABLE III. It is contemplated that these analogs will be useful as TSH antagonists and be useful for the treatment of hyperthyroidism.

It is also within the scope of the invention as illustrated above, that the <u>alpha</u> and <u>beta</u>-subunit sequences in the chimeras can be from any vertebrate (e.g., mammal, bird, amphibian, reptile or fish).

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Further, the invention contemplates modifying, e.g., increasing the stability of glycoprotein hormones in general, and in particular, of the chimeras of the invention (if desired) as by modifying the orientation and/or number and/or positions of the cysteine residues. It is to be noted that notwithstanding the various structural changes discussed herein, the molecules of the invention fold into the proper conformation as is apparent from the fact that they have biological activity.

#### OTHER ANALOGS

The invention also provides for other 20 glycoprotein hormone analogs wherein methionine and other amino acids have been substituted. Such analogs are shown in TABLES VII and VIII. The structures of the various native glycoprotein hormones are such that they have resisted enzymatic digestion that would permit 25 identification of their disulfide bonds (1, 2). contemplated that constructs such as those shown TABLE VIII, which have been modified to introduce additional sites for cleaving the protein, will be useful for determining the disulfide bonds of the hormones. 30 These particular analogs (containing additional methionine residues) would be cleaved with cyanogen bromide (CNBr) and the sequences of the resulting peptides determined. Based on these sequences, possible to identify which Cys residues from particular disulfide-linked pairs. The method can also be used to 35 show that mutagenesis has not altered the "normal" disulfide pattern of analogs such as those described herein, and is another method for examining folding of the protein.

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#### METHODS OF USE

invention The compounds of the can formulated into suitable pharmaceutical, physiologically compositions. These compositions active intravenously, intramuscularly, administered parenterally, orally or by other routes appropriate for The amount of the active analog of the proteins. invention will be used in an amount sufficient to promote the desired physiological response but preferably not in an amount in excess of that necessary to cause an optimum response. The amount may vary from a few nanograms to about 10 kilograms, the amount being dependent of the human or animal to whom the administration is made, the condition to be treated and other variables to be taken One skilled in the art will determine into account. without undue experimentation the optimum amounts to be The compositions will normally include a administered. pharmaceutically or physiologically acceptable carrier or diluent with the active protein of the invention. compositions may be in any suitable form, like injectable preparations.

vaccines of the present incorporating the compounds of the invention, can be prepared according to known methods where the antigen is combined in admixture with a suitable vehicle. vehicles include, for example, saline solutions, various known adjuvants, or other additives recognized in the art for use in vaccine compositions. It is also contemplated that recombinant cells or organisms, live, attenuated or be utilized to deliver antigen(s) might Such vaccines will contain an effective vaccination. amount of the glycoprotein of the invention and a suitable amount of vehicle in order to prepare a vaccine. useful for effective administration to the host. For antigens, further background details on preparation, see "Antibodies: development and

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Laboratory Manual", E. Harlow and D. Lane eds., Cold Spring Harbor Laboratory Press, Cold Spring Harbor, 1988; "Advances in Veterinary Science and Comparative Medicine, vol. 33, Vaccine Biotechnology", J.L. Bittle and F.A. Murphy eds., Academic Press, San Diego, 1989 and "New Trends and Developments in Vaccines," A. Volles and H. Friedman, eds., University Park Press, Baltimore, 1978, which are hereby incorporated by reference.

10 EXAMPLES

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The following EXAMPLES are provided by way of illustration and not by way of limitation. One skilled in the art is capable without undue experimentation, to 15 vary the components and thus construct other analogs. EXAMPLE 1 describes the insertion of hCG alpha-subunit and beta-subunit cDNAs into the eukaryotic expression vector pSVL. EXAMPLE 2 describes the construction of the pKBM, pKBM-hCG-alpha and pKBM-hCG-beta. 20 EXAMPLE 3 describes the construction of pKBM-hCG-beta', which contains two silent mutations in the hCG-beta cDNA nucleotide sequence resulting in the removal restriction sites. EXAMPLE 4 describes the cassette mutagenesis to construct analog F8, a deletion 25 construct (see TABLE VII). EXAMPLE 5 describes a chimeric hCG/hFSH beta-subunit, analog B9 (see TABLE II). EXAMPLE 6 describes the construction of a chimeric hCG/hFSH <u>beta-subunit</u>, analog B11 (see TABLE II). EXAMPLE 7 describes the construction of analogs H3 and 30 H6, chimeric human/bovine alpha-subunits (See TABLE IX). EXAMPLE 8 describes the construction of the expression vector pCM, which can be used to stably transfect COS-1 and COS-7 cells. EXAMPLE 9 describes the construction of pBMT2X-hCG-alpha, pBMT2X-hCG-<u>beta</u> and 35 expression constructs for use in C127 cells. EXAMPLE 10 describes techniques used in mutagenesis and preparation analogs. EXAMPLE 11 describes transfection mammalian cells(COS-1, COS-7, C127, CHO) with plasmid EXAMPLE 12 describes immunological assays of the

analogs. EXAMPLE 13 describes assays of receptor binding. EXAMPLE 14 describes assays of receptor stimulation (steroidogenesis or cAMP accumulation).

Unless indicated otherwise, standard molecular 5 41) were techniques (e.q., ref. biological Restriction endonucleases and other DNA modification enzymes (e.g., T4 DNA polymerase, Thermus aquaticus (Taq) DNA polymerase, T4 DNA ligase, calf intestine alkaline alkaline phosphatase) phosphatase, bacterial 10 Biolabs, Inc., Bethesda England obtained from New Laboratories (Life Technologies), Research (BRL), Boehringer Mannheim, United States Biochemical Corp., or Cetus, and were used as suggested by the manufacturer unless indicated otherwise. 15 efficiency competent HB101 and DH5-alpha E.coli were obtained from BRL and transformed as suggested by the manufacturer with modifications as noted. DNA sequencing was performed using a Sequenase (TM) kit (United States Biochemical Corp.) as suggested by the manufacturer. 20 (COS-1 or COS-7), C 127 and CHO-K1 (CHO) cells were obtained from the American Type Culture Collection (ATCC-COS-7 and C127cells were Rockville, MD). cos-1, in Dulbecco's modified Eagle's routinely maintained medium (DMEM) containing 10% fetal calf serum (FCS) or 25 10% supplemented calf serum. CHO cells were cultured in F-12 medium containing 10% FCS as suggested by ATCC. reagents were obtained GIBCO from Tissue culture HyClone (Life Technologies) Inc. and Laboratories Monoclonal antibodies were provided Laboratories, Inc. 30 by Drs. R.E. Canfield (Columbia Univ., New York NY), R.J. Ryan (Mayo Clinic, Rochester, MN), E.G. Armstrong and R. Wolfert (Hybritech Corp., San Diego, CA), and R. Krogsrud and S.Berube (Biomega Diagnostics, Montreal, Canada).

#### EXAMPLE 1

# Construction of pSVL-hCG-beta and pSVL-hCG-alpha

The cDNAs for the hCG-alpha and beta-subunits were cloned from human placenta, inserted into the E. coli replicating plasmid pBR322, and sequenced by Fiddes and Goodman (40,43). These clones (pBR322-c-alpha-hCG and pBR322-c-beta-hCG) were obtained from J. Fiddes (California Biotechnology, Mountainview CA) and the cDNA inserts were transferred into the E. coli replicating plasmid/eukaryotic expression vector pSVL (Pharmacia) to create pSVL-hCG-alpha and pSVL-hCG-beta.

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15 FIGURE 1 depicts the nucleotide sequence of the hCG beta-subunit cDNA, which includes the coding sequence for the mature protein and signal sequence, as well as adjoining 5′ and 3′ non-translated regions poly(A)/poly(T) region corresponding to the poly(A) tail 20 of the mRNA. The hCG beta-subunit amino acid sequence encoded by the cDNA is depicted using the standard single-letter code (see legend for TABLE I) under the nucleotide sequence. The numbering scheme is the same as used by Fiddes and Goodman (40).Several restriction 25 sites are shown above the nucleotide sequence. Sites unique in pSVL-hCG-beta or pKBM-hCG-beta (see EXAMPLE 2) are underlined.

Ten micrograms (ug) of plasmid pBR322-c-beta-30 hCG was digested with the restriction enzyme HindIII and the 3'-recessed ends of the cut DNA were filled in with DNA polymerase. The DNA fragment containing the entire 579 base pair (bp) hCG-beta cDNA was purified by electrophoresis in a 1% low melting temperature agarose 35 (Sea Plaque[TM]-FMC, Inc.)/ethidium gel bromide described elsewhere (42). The DNA fragment containing the hCG-beta cDNA was visualized on a UV light box and recovered by isolating a gel slice containing the DNA with a razor blade. Ten micrograms of plasmid pSVL were

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cut with the restriction enzyme SmaI and the 5'-phosphate groups were removed from the ends of the cut DNA by treatment with calf intestine alkaline phosphatase. (4889 bp) was purified linearized vector DNA electrophoresis and isolated as described above. The gel slices containing the hCG-beta cDNA and linearized pSVL 65° C. melted by heating at DNA were vector 10 minutes, after which 8 microliters of the hCG-beta cDNA were combined with 2 microliters of pSVL and ligated overnight at 15° C. with 400 units T4 DNA ligase (New England Biolabs) in a volume of 20 microliters. ligation mix was the heated at 65° C. for 10 minutes, diluted with 180 microliters of CaCl2/PIPES buffer (60 mM CaCl2, 10 mM piperazine-N-N'-bis[2-ethanesulfonic acid], pH 7), and 30 microliters of the diluted reaction was used to transform E. coli DH5-alpha cells. Plasmid DNA from ampicillin-resistant clones were screened by codigestion with the restriction enzymes PstI and XbaI, which produces a DNA fragment of approximately 233 bp if the cDNA is inserted in the correct orientation (i.e., proper orientation for transcription of sense mRNA).

FIGURE 2 depicts the nucleotide sequence of the hCG alpha-subunit CDNA, which includes the sequence for the mature protein and signal sequence, as well as adjoining 5' and 3' non-translated regions. The hCG alpha-subunit amino acid sequence encoded by the cDNA is depicted using the standard single-letter code under the nucleotide sequence. The numbering scheme is the same as that used by Fiddes and Goodman (43). the nucleotide restriction sites are shown above Sites unique in pSVL-hCG-alpha or pKBM-hCGsequence. alpha (see EXAMPLE 2) are underlined. The cDNA for the hCG alpha-subunit (43) was placed into pSVL in the same manner as described for hCG-beta, except that screening for pSVL-hCG-alpha was performed by digestion with XbaI alone and selection of clones exhibiting an approximately 400 bp fragment.

These vectors were used to express hCG subunits together or separately in COS cells (44) and as parent constructs for developing plasmids to express and/or modify glycoprotein hormone subunits. In addition, many of the modifications to the cDNA sequences were made directly in pSVL-hCG-beta and pSVL-hCG-alpha. Other plasmid vectors used in the course of this work are pTZ19R (46), pUCX2 (described below), pBMT2X (47) and pCM (described below).

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#### EXAMPLE 2

Construction of pKBMt, pKBM, pKBM-hCG-alpha, pKBM-hCG-beta, pUCX2

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The <u>E. coli</u> replicating plasmid pKBM is a derivative of pUC18 (45) which has been modified to contain the following polylinker, (starting with the HindIII site of pUC18), HindIII, XhoI, XbaI, SacI (SstI), BamHI, BssHII, SalI, SacII (SstII), and EcoRI. This vector was developed to facilitate cassette mutagenesis of glycoprotein hormones. Unlike pSVL, pKBM and pKBMt lack cleavage sites for the restriction enzymes PstI, PpuMI, NaeI and BsmI among others. The absence of these sites in pKBM facilitated cassette mutagenesis using these sites in hCG <u>alpha</u> and <u>beta-subunit cDNAs</u>.

FIGURE 3 depicts the oligodeoxynucleotide cassettes used in the construction of pKBMt (3a) and pKBM (3b). Restriction endonuclease cleavage sites encoded by each cassette are indicated.

Two micrograms of pUC18 were digested with the restriction endonucleases HindIII and XbaI and purified by electrophoresis in low melting point agarose as described in EXAMPLE 1. The two synthetic DNA oligonucleotides depicted in FIGURE 3a were synthesized using an Applied Biosystems 380B DNA Synthesizer and the phosphoramidite method (51), purified by polyacrylamide

gel electrophoresis (as described in Applied Biosystems User Bulletin No. 13, April 1, 1987), and quantified by spectrophotometry (absorbance The 260 nm). at combining then annealed by were oligonucleotides 1 microgram of each oligonucleotide in 100 mM KCl, 10 mM 5 Tris pH 8.0, 1 mM EDTA (total volume=200 microliters) 100° C. for five minutes and cooling for heating at **Approximately** room temperature. at 30 nanograms of HindIII/XbaI cut pUC18 was combined with of the annealed oligonucleotides 100 nanograms 10 ligated overnight at 15° C. with 1 unit T4 DNA ligase in a 20 microliter Research Laboratories) (Bethesda The ligation reaction was heated at 650 C. for of 180 microliters with diluted and 10 minutes Thirty microliters of the diluted ligation CaCl<sub>2</sub>/PIPES. 15 reaction was used to transform E. coli K-12 strain JM83 (87) which were then plated with 50 microliters of a 2% 5-bromo, 4-chloro, 2-indoly1-beta-D-galactoside Plasmid DNA from ampicillin-resistant DMSO solution. blue colonies were screened with XhoI to verify the 20 presence of the cassette. The resulting plasmid was To construct the plasmid pKBM, two designated pKBMt. synthesized, oligonucleotides were DNA additional annealed to form the cassette depicted in FIGURE 3b, and ligated with EcoRI/BamHI digested pKBMt. Plasmid DNA 25 from ampicillin-resistant blue colonies were screened with BssHII, and plasmids containing the new cassette The hCG alpha and beta-subunit were designated pKBM. cDNAs were subsequently transferred from pSVL into pKBM between the XhoI and BamHI sites to create pKBM-hCG-alpha 30 is another The vector pUCX2 pKBM-hCG-beta. derivative of pUC18 with an altered polylinker. polylinker in pUCX2 is (starting from the former HindIII site of pUC18), BssHII, SacII, HindIII, XhoI, and EcoRI.

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#### EXAMPLE 3

# Construction of pKBM-hCG-beta'

An analog of the hCG <u>beta</u>-subunit cDNA, called hCG-<u>beta</u>', containing two silent mutations which eliminate restriction sites has been the starting construct for many mutants.

FIGURE 4a depicts the nucleotide sequence of 10 the hCG-beta' DNA. As with the hCG-beta cDNA it includes the coding sequence for the mature protein and signal sequence, adjoining 5' and 3' non-translated regions and a poly(A)/poly(T) region corresponding to the poly(A) 15 tail of the mRNA. The encoded amino acid sequence depicted under the nucleotide sequence is unchanged from that of hCG-beta. The locations of the silent mutations in the DNA sequence are indicated by solid triangles above the base pairs at positions 229 and 232. numbering scheme is the same as that used by Fiddes and 20 Goodman (40). Several restriction sites are shown above the nucleotide sequence. Sites unique in pSVL-hCG-beta' or pKBM-hCG-beta' are underlined.

FIGURE 4b depicts the oligonucleotide cassette used to construct pKBM-hCG-beta'. As in FIGURE 4a, the mutated bases are indicated by solid triangles.

Approximately 5 micrograms of pKBM-hCG-beta was 30 digested with the restriction endonucleases PstI OxaNI (OxaNI is an isoschizomer of Bsu36I and MstII) and restriction fragment purified electrophoresis in low melting point agarose as described Two synthetic DNA oligonucleotides (FIGURE 4a) were synthesized, purified and annealed as described in 35 EXAMPLE 2. Two hundred nanograms of PstI/OxaNI cut pKBMhCG-beta was combined with 80 nanograms of the annealed oligonucleotide cassette, ligated and then transform E. coli JM83 as described. Ampicillinresistant white colonies were picked and plasmid DNA from these clones was screened by digestion with the restriction endonuclease PpuMI, which cuts out an approximately 200 bp fragment from pKBM-hCG-beta, but no fragment from pKBM-hCG-beta'. The hCG-beta DNA in pKBM-hCG-beta' differs from the native hCG-beta cDNA in that it contains unique NaeI and PpuMI restriction sites, facilitating its modification by cassette mutagenesis.

10 EXAMPLE 4

## Construction of Analog F8

depicts the FIGURE 5a DNA oligonucleotide cassette used to construct analog F8 and the encoded amino acid sequence. The deletion of residues 47-50 (//) and substitution of proline for the alanine at position 51 (solid triangle) in the native hCG beta-subunit are These oligonucleotides were synthesized, indicated. purified and annealed as described in EXAMPLE 2, and then ligated with PstI/OxaNI digested pKBM-hCG-beta as described in EXAMPLE 3. Plasmid DNA from ampicillinresistant clones of transformed JM83 E. coli was screened for the presence of the insert by digestion with PpuMI as described in EXAMPLE 2. Plasmids with the proper insert are denoted pKBM-F8 (also referred to as pKBM-beta-delta 3 or pKBM-beta-del 47-50, 51P). encoding the analog was transferred in to the expression vector pSVL using the restriction endonucleases XhoI and BamHI as described for B9 in EXAMPLE 5, creating pSVL-F8.

### EXAMPLE 5

## Construction of Analog B9

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FIGURE 5b depicts the DNA oligonucleotide cassette used to construct analog B9 and the encoded amino acid sequence. The substitution of hFSH <u>beta</u>-subunit residues 88-91 (DSDS) for the corresponding

residues of the hCG beta-subunit, i.e., residues 94-97 (RRST) are indicated by solid triangles, and the recognition site for restriction endonuclease ScaI is also noted. Approximately 25 micrograms of plasmid pKBM-5 hCG-beta' was used in a time-course digestion with restriction endonuclease 20 units of ApaLI 100 microliter volume. A portion of the reaction was removed, placed on ice, and adjusted to 20 mM EDTA at time 0 (12 microliters, taken prior to addition 10 10 (12 microliters), enzyme), and after 20 (24 microliters), 30 (24 microliters), 45 (12 microliters) and 60 (16 microliters) incubation at 37° C. These were then purified electrophoresis in low melting point agarose as described 15 in EXAMPLE 1. In this particular time-course optimal linearization of the plasmid was seen after 20 minutes. The bands containing single cut plasmid (approximately 3280 bp) were cut from the gel, transferred to microfuge tube, and the gel slices melted by incubating at 2Ó 65° C. for 10 minutes, after which they were diluted with 200 microliters deionized water. Two hundred microliters of phenol (adjusted as described in ref. 41) were then added, the tube vortexed, and then placed in a dry ice/ethanol bath for 15 minutes, after which they were 25 centrifuged at full speed in an Eppendorf microfuge at for 15 minutes. The aqueous fraction was collected and the organic fraction back extracted with an additional 200 microliters deionized water. The aqueous phase from this second extraction was pooled with the 30 aqueous phase from the initial extraction, and then with 400 microliters of chloroform/isoamyl extracted alcohol (24:1 mixture as described in ref. 40), after which the DNA was recovered from the aqueous phase by precipitation with sodium acetate and ethanol. 35 was resuspended in 14 microliters deionized water. digested with PpuMI, and purified by electrophoresis. oligonucleotides depicted in FIGURE 5b synthesized, purified and annealed, and then ligated with the ApaLI/PpuMI digested pKBM-hCG-beta' DNA. Plasmid DNA

from ampicillin-resistant clones of transformed JM83 E. coli was screened for the presence of the insert by digestion with ScaI, which cuts pKBM-hCG-beta' once and cuts pKBM-hCG-beta' containing the insert twice. vector is denoted herein as pKBM-B9. The coding sequence for analog B9 was subsequently transferred to pSVL by digesting pKBM-B9 with XhoI and BamHI, gel purifying the approximately 600 bp fragment, and ligating XhoI/BamHI digested pSVL. Purified preparations of this vector, pSVL'B9, also referred to as pSVL-beta-D or pSVLbeta-CF94-97, were sequenced and used to co-transfect COS7 cells with pSVL-hCG-alpha.

#### EXAMPLE 6

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#### Construction of Analog B11

In some instances multiple cassettes were ligated into a vector, as for example the assembly of pSVL-B11. This analog, in which amino acids 94-145 in the mature hCG <u>beta</u>-subunit are replaced with amino acids 88-111 from the mature hFSH <u>beta</u>-subunit, was built using plasmid pSVL-B9 (EXAMPLE 6) and two oligonucleotide cassettes.

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FIGURE 5c depicts the DNA oligonucleotide cassettes (A and B comprising oligonucleotides A+ and A-, and B+ and B-, respectively) used to construct analog B11 from B9, and the encoded amino acid sequence. residues beta-subunit 95-111 substitution of hFSH (TVRGLGPSYCSFGEMKE) for the hCG beta-subunit residues 101-145 (GGPKDHPLTCDDPRFQD...Q, see FIGURE 1 for full sequence) are indicated by the solid triangles, recognition sites for restriction endonucleases StuI and PstI are noted.

The cassettes were designed so that they could be ligated together (they overlap by 4 complementary, non-palindromic bases) and into the ScaI and BamHI sites

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of pSVL-B9. The oligonucleotides for the cassettes were synthesized as described for pKBM-hCG-beta' and purified oligonucleotide purification cartridges The 5'- ends of oligonucleotides A-Applied Biosystems. and B+ were phosphorylated using T4 polynucleotide kinase ligation of the two Oligonucleotides A+ and A-, and B+ and B-, were then combined and annealed described as above. Four micrograms of pSVL-B9 DNA was partially digested with restriction endonuclease ScaI using a time-course, purified by gel electrophoresis, and extracted from the gel as described. The purified ScaI linearized pSVL-B9 DNA was then resuspended in 16 microliters deionized water, digested with restriction endonuclease BamHI, purified by gel electrophoresis, and combined with the cassettes in annealed a ligation reaction. Ampicillin-resistant clones containing plasmids with the desired insert were isolated by digesting plasmid DNA with PstI, for which there are two recognition sites in pSVL-B9 and three in plasmids with the desired insert, which are denoted pSVL-B11 (also pSVL-beta-DG or pSVLbeta-CF94~117).

#### EXAMPLE 7

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#### Construction of Analogs H3 and H6

FIGURE 6 depicts the nucleotide sequence of the synthetic bovine <u>alpha</u>-subunit cDNA, constructed from multiple cassettes in pTZ19R, which includes the coding sequence for the mature bovine <u>alpha</u>-subunit protein and the human <u>alpha</u>-subunit signal sequence. The <u>alpha</u>-subunit amino acid sequence encoded by the cDNA is depicted using the standard single-letter code under the nucleotide sequence. Several restriction sites are shown above the nucleotide sequence. Sites unique in pSVL-b-alpha or pTZ19R-b-alpha are underlined. Several restriction sites in common with the native human cDNA

have been used to construct chimeric alpha-subunit analogs.

The XbaI site was used to construct analogs H3 (also referred to as pSVLhum/bov 34) 5 and H6 referred to as pSVLbov/hum 34), which are depicted in TABLE IX. To build analog H3, approximately 5 micrograms of pSVL-b-alpha or pSVL-hCG-alpha were digested with the restriction endonucleases XbaI and EcoRV and purified by 10 gel electrophoresis. The large fragment from the digest of pSVL-hCG-alpha and the small fragment from the digest of pSVL-b-alpha were ligated and used to transform DH5-Plasmid DNA from ampicillin-resistant alpha E. coli. clones was screened by digestion with XbaI (pSVL-H3 has 15 single XbaI site, pSVL-hCG-alpha has two). To build analog H6, approximately 2 micrograms of pSVL-b-alpha was digested with restriction endonucleases XbaI and SalI, purified, the and large fragment Approximately 5 micrograms of pSVL-hCG-alpha was digested with XbaI and SalI, gel purified and the middle fragment 20 The saved fragments of pSVL-b-alpha and pSVL-hCGligated and used to transform DH5-alpha E. coli. Plasmid DNA from ampicillin-resistant clones screened by digestion with XhoI and BamHI, with 25 positive clones of pSVL-H6 exhibiting an approximately 600 bp fragment as opposed to the approximately 380 bp fragment obtained from pSVL-b-alpha.

#### EXAMPLE 8

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#### Construction of pCM, pCM-hCG-alpha, pCM-hCG-beta

The vector pCM contains the Simian Virus 40 of replication, early origin promoter 35 polyadenylation sequence; the neomycin (aminoglycoside) phosphotransferase gene, ampicillin resistance gene, and bacterial origin of replication (i.e., the EcoRI/BamHI fragment) from pMVBneo (47), the SV40 origin replication and late promoter, and synthetic polylinker

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(XhoI-XbaI-SmaI-SstI-BamHI) from pSVL (<u>i.e.</u>, the HindIII/BamHI fragment), and the HindIII/EcoRI fragment of Bovine Papilloma Virus (BPV), which contains BpV genes E6, E7, E8, and a portion of E1, the two plasmid maintenance sequences, and an enhancer region (for review see ref. 48). The HindIII/EcoRI fragment from BPV which we placed into pCM has been shown to be capable of controlling replication of plasmids with an SV40 origin of replication permitting the creation of COS cell lines stably transformed with an episomal vector (49).

The plasmid pCM was constructed as follows. Two micrograms of pSVL was digested with restriction endonucleases EcoRI and BamHI and the smaller fragment (approximately 2000 bp) purified by electrophoresis as in EXAMPLE 1. Two micrograms of plasmid pMVBneo (47), obtained from G.N.Pavlakis (National Cancer Institute, Frederick, MD), was digested with EcoRI and BamHI and the large fragment (approximately 5000 bp) gel These were then ligated and plasmid DNA from ampicillin-resistant clones of DH5-alpha E. coli screened by digestion with HindIII. Clones exhibiting HindIII cut fragments of approximately 1100, 2950 and 3000 bp were positive for the intermediate plasmid, designated Additional DNA from these clones was then pSVLneo. screened by co-digestion with either EcoRI and BamHI (which was expected to yield fragments of approximately 2000 and 5000 bp) or EcoRI, BamHI and HindIII (which was expected to yield fragments of approximately 450, 500, 1100, 2500 and 2500bp). To assemble the final vector, pCM, approximately 5 micrograms of DNA from a clone of pSVLneo was partially digested with 2 units of HindIII in a reaction from which samples were removed, adjusted to 25 mM EDTA (to inactivate the enzyme) and placed on ice at the following time points; 0 (10 microliters), 10 (22 microliters), 20 (22 microliters), 30 (22 microliters) and 45 (22 microliters) minutes after addition of enzyme. These samples were gel purified and the linearized plasmid (approximately 7000 bp) was cut

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from the gel and transferred to a microfuge tube. slice was melted by incubation at 65°C. for 10 minutes, and 10 microliters was transferred from the microfuge tube to a second tube containing 6 microliters of water, 2 microliters React3 (TM) (BRL) and 20 units of This tube was incubated at 37° C. for an hour, and the approximately 6500 bp EcoRI/HindIII cut band purified by gel electrophoresis. The plasmid pBMT2X (47), obtained from G.N. Pavlakis, was digested with HindIII and EcoRI, gel purified, and the 2200 bp fragment corresponding to the 2200 bp fragment of BPV cut from the ligated with the This fragment was EcoRI/HindIII cut fragment of pSVLneo, and plasmid DNA from ampicillin-resistant clones of DH5-alpha E. coli digested with HindIII to identify plasmids with the 2200 insert from pBMT2X (positive clones exhibit bands Fof approximately 1100, 3000 and 4800 bp). The new plasmid The hCG-beta cDNA was was designated pCM or pSVLneoBPV. inserted between the XhoI and BamHI sites of pCM and plasmid DNA from ampicillin-resistant clones was screened by digestion with PstI or XhoI and BamHI. This new plasmid was designated pCM-hCG-beta. The hCG-alpha cDNA and DNAs encoding hormone analogs were placed into pCM-in similar fashion.

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The new vector pCM combines attributes of the two principal existing types of eukaryotic expression systems, i.e., transient high level expression and stable expression, into a single system. When transfected into COS cells using procedures described in EXAMPLE 11, pCMhCG-beta provided rapid, high level expression recombinant proteins, superior to that obtained with When these transfected COS cells are pSVL-hCG-beta. incubated with medium containing 500 micrograms/ml of synthetic aminoglycoside G418 (Geneticin-GIBCO) it was possible to establish stably transfected cells which produced recombinant proteins at levels of hundred nanograms/million cells/day for prolonged periods of time (e.g., >30 cell passages), thereby facilitating

the cloning and characterization of gene products. These cells were also successfully frozen, stored in liquid nitrogen for several months, and then recultured without loss of the ability to secrete hCG <u>beta</u>-subunit.

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It was also possible to recover the DNA vector the transformed COS cells and reclone Transformed cells were scraped from bacteria. surface of a culture dish and transferred to a 15 ml The cells were then washed twice with 5 ml cold phosphate buffered saline, resuspended in 400 microliters sodium-dodecyl sulfate (SDS)/10 mM EDTA, and incubated for transferred to a microfuge tube, 20 minutes at room temperature. This mixture was then adjusted to 1M NaCl (by addition of 100 microliters of 5M NaCl) and incubated on ice for at least five hours. After this the cellular debris were centrifuged out of solution (full speed in a microfuge for 5 minutes) and the resulting supernatant containing the plasmid was extracted with phenol and precipitated with ethanol and 10 micrograms of mussel glycogen. The isolated DNA was resuspended in 20 microliters of water. One microliter of this was used to transform DH5-alpha E. coli, yielding 63 ampicillin-resistant colonies (we found that these bacteria can also be selected with G418 due to the presence of the neomycin phosphotransferase gene in the vector, even though it is downstream of the SV40 early promoter). Plasmid DNA was prepared from 24 of these clones and digested with XhoI and BamHI. Twenty three of the recovered plasmids were found to contain a fragment of approximately 600 bp corresponding to the hCG-beta cDNA, whereas one clone did not grow in liquid culture. None of the plasmids gave any evidence of rearrangement. strategy and procedure, previously used transiently transfected COS cells, is now extended to stable transformed cells containing a hybrid BPV-SV40 Vectors of this type could conceivably be used in any cell type supporting an SV40 origin of replication.

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Other related viral origins (e.g., polyoma virus) may work in this system as well. Other promoters, enhancers, selectable markers could also be used, singly or in combination. Vectors such as pCM can be used for genes, characterization of gene modification of cells, and application any other involving selection of particular transfected cells from a pool of cells. Among the analogs that have been subcloned into pCM and expressed in COS cells is analog J2, in which the hCG beta-subunit is anchored to the cell surface via fusion of the transmembrane domain from the Vesicular Stomatitis Virus G-protein (VSV-G) onto the carboxy end of hCG-beta at the SmaI site.

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TABLE XV illustrates the retention of hCG betasubunits on COS-7 cells transformed with pCM-beta-J2 and maintained in 500 micrograms/ml G418. COS-7 transformed with pCM-hCG-beta do not retain much betasubunit on their surface. In this assay confluent 75 cm<sup>2</sup> flasks of cells were cultured for about 30 hours, with serum free DMEM containing 500 micrograms/ml G418. The medium was then removed and the cells were gently washed with 10 ml phosphate-buffered saline (PBS). The cells were then removed from the surface of the flask by incubation for 2 hours at 37° C. with PBS containing 0.1% sodium azide/10 mM EDTA. The cells were resuspended in this and 450,000 cells were added to microfuge tubes containing 75,000 cpm 125I-labelled antibody B105, and if indicated, 300 nanograms unlabeled B105. These were incubated at 37° C. for an hour, diluted with one ml of a 0.9% saline/0.1% bovine serum albumin solution, and the cells were pelleted by centrifuging for 8 minutes at full speed in a microfuge. The supernatant was aspirated and the pellets counted in a gamma counter.

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#### EXAMPLE 9

Construction of pBMT2X-hCG-<u>alpha</u>, pBMT2X-hCG-<u>beta</u>, pBMT2X-F8

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The vector pBMT2X permits stable expression of recombinant proteins in C127 cells (47). hCG subunits and analogs were sub-cloned into the unique XhoI site of (provided by Dr.G.N. Pavlakis, see EXAMPLE 8). Approximately 10 micrograms of plasmids pKBM-hCG and pKBM-hCG-beta were digested restriction endonucleases XhoI and SalI and the approximately 600 bp fragments containing the alphasubunit and beta-subunit cDNAs were isolated as described in EXAMPLE 1. Two micrograms of plasmid pBMT2X was digested with XhoI, the 5' terminal phosphates removed from the DNA with calf intestinal alkaline phosphatase, and the linearized DNA gel purified as described in The XhoI digested pBMT2X and cDNA fragments example 1. were ligated with T4 DNA ligase and used to transform HB101 E. coli as described for DH5-alpha E. coli in EXAMPLE 1. Plasmid DNA from ampicillin-resistant clones digested with BssHII, which linearized containing the cDNA inserts from pKBM. The orientation of the inserts was determined by digestion of the plasmid DNA with XhoI and SstII, with inserts in the proper orientation exhibiting a fragment of approximately 1400 bp. These new plasmids were designated pBMT2X-hCGalpha, pBMT2X-hCG-beta. Plasmid preparations of pBMT2XhCG-alpha and pBMT2X-hCG-beta were subsequently used to transform C127 cells as previously described (50, see also EXAMPLE 11) and cells producing either hCG dimer, or alpha or beta were sub-cloned in the presence of 20-50 uM cadmium chloride (CdCl2). Production of protein by these cells was on the order of 5 micrograms hCG dimer/million cells/24 hours. DNA encoding analog F8 was subcloned pKBM-F8 into pBMT2X in similar fashion and subsequently co-transfected with pBMT2X-hCG-alpha C127 cells and CdCl2 resistant cells subcloned.

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#### EXAMPLE 10

### METHODS OF MUTAGENESIS Oligonucleotide Cassette Mutagenesis

Many constructs have been prepared by DNA oligonucleotide cassette mutagenesis similar to the manner in which pKBMhCG-beta', pKBM-F8, pKBM-B9 or pSVL-B18 were built.

#### Restriction Fragment Mutagenesis

Vectors containing native cDNAs, genes or analogs thereof with restriction sites in common can be used to create new analogs by replacing restriction fragments, as was done in the construction of analogs H3 and H6.

#### Restriction Site Fill-in Insertion Mutagenesis

The analog G7 (also referred to as hCG-beta-55iH or hCG-delta 1) was constructed by cutting pSVL-hCG-beta with OxaNI, filling in the ends of the linearized DNA with T4 DNA Polymerase, religating the vector with T4 DNA ligase, transforming DH5-alpha E.coli, and selecting clones lacking an OxaNI site. Similar procedures can be used to mutagenize other restriction sites with a three base overlap. In addition, cassettes can be used to introduce deletions or insertions.

#### Site-directed Oligonucleotide Mutagenesis

A series of hCG/hLH <u>beta</u>-subunit chimeras were prepared by oligonucleotide site-directed mutagenesis of the hCG-<u>beta</u> cDNA. The construction of one of these constructs, pTZ19R-A4 (also referred to as pTZ19RhCG/hLH96) is briefly described here. Synthetic DNA oligonucleotides were prepared which contained

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regions of DNA encoding hLH amino acids 96-117 flanked by regions homologous to the hCG-beta cDNA sequence. The hCG-beta cDNA was sub-cloned into pTZ19R between the SacI and SalI sites. The various oligonucleotides were annealed to single stranded vector and used to prime synthesis of double-stranded vector with the Klenow fragment of <u>E. coli</u> polymerase I. The resulting plasmids were used to transform bacteria and colonies containing plasmids with the modified cDNA isolated.

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#### Polymerase Chain Reaction Mutagenesis

Several analogs of hCG-beta have been produced using synthetic oligonucleotides and the polymerase chain reaction (PCR). For example, to produce F1, an analog in which the hCG-beta amino acid sequence was truncated at residue 111 (i.e., Cys-110 is the last amino acid) the oligonucleotide synthesized, following was 5'-AAGGACCACCCCTTGACCTGTTAGGATCCTATATA-3', and used as a primer in a PCR reaction with a primer to pSVL which binds 5' to the hCG-beta cDNA. The resulting PCR product cut with XhoI and BamHI, isolated electrophoresis, and cloned into the XhoI and BamHI sites of pSVL using the techniques described previously.

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#### EXAMPLE 11

#### Introduction of Genetic Vectors into Mammalian Cells

30 Both calcium phosphate and DEAE-dextran methods have been used to transfect COS (COS-1 or COS-7) cells with vectors containing glycoprotein hormone subunits or analogs. The calcium phosphate technique modification of a published protocol(50). COS cells to 35 be transfected were plated the day before on 6-cm or 10cm diameter tissue culture plates at a density of 10,000 cells per square centimeter. Plasmid DNA was purified in an ultracentrifuge over a CsCl gradient and sterilized by precipitation with ethanol. The quantity of DNA used in

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the transfection procedure varied from 1 to 20 micrograms per plate of cells. The final volume of the DNA solution was either 400 microliters (6-cm plate) or 1 ml (10-cm Salmon sperm DNA was sometimes used as a carrier. After resuspending the DNA in sterile water, an appropriate volume of 2 M CaCl<sub>2</sub> was added to yield a final concentration of 125 mM of CaCl2 in 200 microliters (for each 6-cm plate) or 500 microliters (for each 10-cm plate). An equal volume of 2X BES-buffered saline (50 mM of N-,N-Bis[2-hydroxyethyl]-2-aminoethanesulfonic acid, 280 mM NaCl, 1.5 mM Na2HPO4; adjusted to pH 6.95) was then added to the DNA/CaCl<sub>2</sub> solution. This mixture was vortexed, incubated at room temperature for 10-20 minutes and then added drop-wise to the plates of cells. cells and DNA were incubated overnight (18-30 hours in a 37° C. incubator under 2.5-6% carbon dioxide. was removed, the cells washed with 1X PBS and then the cells were cultured in complete medium. 24 hours this media was removed, the cells were washed twice with 10 ml PBS, and serum-free DMEM was added. media was harvested 70-80 hours later for recovery of secreted protein.

Transfections using DEAE-dextran were done published protocol (86). according to a procedure, one million cells were plated on 100 mm plates the day before the transfection. Plasmid as described above, then sterilized prepared and sterile water, adjusted to 1x **PBS** resuspended in 200 micrograms/ml **DEAE-dextran** concentration and Media was aspirated from the plates of cells, 1.2 ml. and the plates were washed twice with PBS. The DNA/DEAEdextran solution was then added to the plates, which were placed in a 37° C. incubator for 40 minutes. After this time, 10 ml DMEM made 100 uM with chloroquine was added to each plate. The plates were incubated for 3-4 hours, after which the media was replaced with medium containing 10% DMSO. Following 2-3 minutes at 37° C. the medium was aspirated, the plates were washed with PBS, covered with

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10 ml DMEM containing 10% FCS or serum substitute (Calf Supreme or Calf Serum, Supplemented from GIBCO) and placed back in the incubator. After 12-24 hours the medium was removed, the plates washed twice with 10 ml PBS, and then covered with 6 ml serum-free DMEM. The medium was harvested 70-80 hours later for recovery of secreted protein.

Stable transfections of COS, C127 or CHO cells were performed using the calcium phosphate procedure Stable clones of C127 cells transfected described above. with pBMT2X-based plasmids were selected by resistance to CdCl<sub>2</sub> (10-80 mM) starting two days after transfection. Cells transfected with plasmids containing the gene for neomycin (aminoglycoside) phosphotransferase (e.g, pcMplasmids) were selected with the synthetic aminoglycoside G418 (Sigma) by adding 500 micrograms/ml G418/ml tissue culture medium (COS) 1000 micrograms G418/ml medium(CHO) starting 1-4 after transfection. After this the cells were routinely maintained under selection pressure.

#### EXAMPLE 12

#### 25 Sandwich Assays

Immunological properties of the hormones and analogs were determined using monoclonal sandwich assays essentially as previously described by Moyle et al. (5). These assays were used to determine whether the analogs folded similar to the native molecule and formed alpha, beta-heterodimers which were secreted into the culture medium.

TABLE XIII summarizes results of the immunological characterization of hCG/hFSH chimeras. All assays employed B105 as "capture" antibody, except for assays of hFSH, which used B601, B602, A113 or E501 (see below). The various labeled detection antibodies are

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The specificity of these in the TABLE. antibodies has been described previously (5-7, 52, 53). In these assays, 1 microgram of capture antibody (e.g., B105) was adsorbed to the surface of individual wells in 96-well microtiter plates by adding 50 microliters of a 0.9% NaCl/20 micrograms/ml antibody solution (made with distilled water) to each well and incubating at 37° C. Unbound antibody was removed and nonfor 1 hour. specific sites were blocked with bovine serum albumin (BSA) by filling each well with a 0.1% BSA/0.9% NaCl solution and incubating at 37° C. for 1 hour. standards or analogs were added to each well (volume 50 microliters) incubated for 1 hour at adiusted to 37° C. to permit the analogs to bind to the adsorbed antibody. The wells were washed with distilled water to remove unbound analogs and 50,000 counts per minute (cpm) "detection" antibody in BSA/saline labeled added to the well and (50 microliters volume) was incubated at 37° C. for 1 hour. Next, the unbound antibody was removed by aspiration, the plates washed with deionized water, and the individual wells cuts out and counted in a gamma counter. Detection antibodies were radioiodinated by the iodogen method as previously described by Moyle et al. (54).

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ability of TABLE XIII illustrates the monoclonal antibody sandwiches which detect the alpha, beta-heterodimer (B105 capture, A113 detection), beta-subunit or hCG (B105 capture, B101 detection), and analogs with <a href="https://example.com">beta</a>-subunit epitopes from both hCG (B105) The ability of the radioiodinated and hFSH (B601). detection antibody to bind to the captured hormone or analog relative to hCG (except for B601) is indicated in Note that the data in TABLE XIII TABLES XIII. expressed as either ++ (cpm bound = 50-120% of those bound to a urinary hCG standard), + (cpm bound = 10-49% of those bound to the standard), or - (<10%binding), and nt indicates not tested. For B601, a monoclonal antibody specific for hFSH, binding to the various analogs is

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described relative to their reactivity in an hFSH assay, i.e., E501/B601 sandwich assay. In this assay, employing E501 capture, B601 was found to bind chimeras containing hFSH residues 33-52 indistinguishably from recombinant hFSH. The immunologic data demonstrates that the analogs fold and bind to the alpha-subunit. It also demonstrates that structural features of one hormone (e.g., the B601 epitope in FSH) can be transferred to another hormone, even in regions of very low homology.

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TABLE XI illustrates monoclonal antibody binding to alpha-subunit analogs co-expressed with hCG beta-subunit. The ability of the radioiodinated detection antibody to bind to the captured hormone or analog relative to hCG is expressed as either ++++(cpm bound = >200% of those bound to a urinary hCG standard), +++(cpm bound = >110% and <200% of those bound to a urinary hCG standard), ++ (cpm bound = 50-110% of those bound to a urinary hCG standard), + (cpm bound = 10-49% of those bound to the standard), or - (<10% binding), and nt indicates not tested. These assays employed B105 as capture antibody. All of these analogs form the alpha, beta-heterodimer (B105 capture, B109 Although antibody B109 is directed against an epitope on the hCG beta-subunit, it binds much better to the alpha, beta-heterodimer than the free beta-subunit, which indicates that it recognizes a conformational change in hCG-beta caused by combination of beta-subunit with alpha-subunit. The data indicate that all of the alphasubunit analogs are capable of inducing this conformational change in hCG-beta. The data illustrate that none of the antibodies directed against the hCG-alpha subunit (A101, A102, A109, A112, A202 and A501) bind well to the bovine alpha-subunit. Replacement of bovine amino acid sequence with sequences from the human alpha-subunit enables these antibodies to bind. some instances, replacement of a discrete contiguous sequence is sufficient to obtain binding similar to that seen with hCG (e.g., A102, A112, A202).

antibodies, a larger contiguous region (A101) or several separate regions (A109, A501) must be substituted to develop antibody binding to the bovine/human chimera. (See TABLE IX for analog sequences) These binding data are summarized in FIGURE 7, which relates binding of hCG alpha-subunit antibodies and the amino acid sequence of the alpha-subunit.

#### EXAMPLE 13

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#### Assay of Receptor Binding

Two types of assays were used to characterize receptor binding. One is a competitive assay while the other is a sandwich or BioIRMA assay. Both have been described previously by Moyle et al. (20, 54).

Membrane bound LH receptors were obtained from rat corpora lutea super-ovulated with pregnant mares serum gonadotropin and hCG (54). Membrane bound FSH receptors were obtained from bovine calf testes (11). In the BioIRMA, hormones or analogs were incubated with a preparation of membrane bound receptors. Hormone which became bound to the receptors was quantified using radiolabeled B105 monoclonal antibody (6) which has high affinity for an epitope on the hCG beta-subunit that remains exposed after the hormone has complexed with receptors.

TABLE XIV illustrates the ability of <u>alpha</u>-subunit analogs co-expressed with hCG <u>beta</u>-subunit to bind LH receptors.

Further characterization of the analog's receptor binding properties was determined by radioligand receptor assays as described by Moyle et al. (20). This assay is a competitive displacement assay in which the analogs compete for receptor binding with radioiodinated hCG and hFSH.

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FIGURE 9 illustrates tha abilities of hCG, hFSH and analogs to inhibit the binding of \$^{125}I-hCG\$ to LH receptors. Their ability to inhibit \$^{125}I-hCG\$ binding correlates to their ability to bind LH receptors. Data from each analog are marked using the nomenclature in TABLE II. Data from analogs with hCG-like activity are plotted with filled boxes, rhCG (---), B15 (---), B27 ('''); data from analogs with hFSH-like activity are plotted with open boxes, rhFSH (-), B11 (---), B17 (----), B18 (---); and data from analogs that behave differently

from both hCG and hFSH are plotted with open triangles,

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15 FIGURE 10 illustrates the abilities of hCG, hFSH and analogs to inhibit binding of <sup>125</sup>I-hFSH to FSH receptors. Their ability to inhibit <sup>125</sup>I-hFSH binding correlates to their ability to bind FSH receptors. Representation of hormones and analogs is the same as for LH receptor binding in FIGURE 9.

B9 (--) and B16 (---).

FIGURE 13 illustrates the ability of the <u>alpha</u>, <u>beta-F8</u> heterodimer to inhibit <sup>125</sup>I-hCG binding to LH receptors. These data show that this analog binds to LH receptors nearly as well as hCG.

For the competitive displacement assays the recombinant hormones and analogs were concentrated using Amicon Centriprep-10 concentrators and quantified by dimer-specific sandwich assays employing monoclonal antibodies to human alpha-subunit (A110, A113) and hCG beta-subunit (B105, B108) using the procedure described above. Urinary derived hCG (CR121 or CR123) was used as the reference standard. Similar results were obtained using either alpha or beta-subunit capture, and with different pairs of antibodies. In addition, analogs that bound anti-hFSH-beta antibody B601 gave comparable results when quantified in an hFSH dimer-specific sandwich assay. Inhibition of labeled hCG binding to LH

receptors from luteinized rat ovaries was determined as The FSH radioligand-receptor assays were described. conducted essentially as previously described except that the binding incubation was carried out at 37° C. Bovine testes membranes used in the FSH receptor assays were a gift from Dr.D.J. Bolt (Beltsville Research Center, Beltsville, Maryland). Agricultural Highly purified urinary hCG (CR121 or CR123) pituitary hFSH (NIADDK-hFSH-I-3 and AFP-4822-B) were used as reference standards and for preparation of label using the iodogen-based procedure. Averaged results from triplicate tubes are presented graphically as percent displacement of total 125I-hCG or 125I-hFSH binding measured in the absence of inhibitor.

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#### EXAMPLE 14

#### Assay of Hormone Responses

20 The ability of hCG, FSH, and the analogs to stimulate steroidogenesis was monitored by radioimmunoassay of testosterone production by rat Leydig cells and estradiol synthesis by rat granulosa cells using procedures similar to those described previously (20, 55).

Rat Leydig cells were isolated and prepared for incubation similar to the procedure described previously Briefly, this procedure consisted of digesting decapsulated rat testes from 22- to 40-day-old Sprague-Dawley rats (Charles River, Wilmington, Mass.), in Krebs Ringer buffer (58) containing 1 mg HEPES in place of (Worthington, and 1 mg collagenase/ml bicarbonate Freehold, N.J.) until the tubules dissociated from one another (10 to 20 minutes). After the addition of 25 ml of enzyme-free buffer to the digestion flasks, the tubules were allowed to sediment and the interstitial cells were decanted into centrifuge tubes. Following centrifugation at 70 x g for 10 minutes the cells were

resuspended in the buffer containing 1 mg of bovine serum albumin and counted in a hemocytometer. The routine steroidogenesis assay consisted of adding 50 microliters of the cell suspension containing approximately 350,000 nucleated cells to 50 microliters of hormone or analog in a plastic tube (12 x 75 mm), and measuring the testosterone produced after a 3-hour incubation at  $37^{\circ}$  C.

Testosterone was measured with 10 radioimmunoassay procedure utilizing antisera obtained from Cambridge Medical Diagnostics (Billerica, MA). this procedure, an aliquot of unextracted incubation medium (50 microliters) containing approximately 175,000 added cells was to 0.2 pmol of <sup>3</sup>H-testosterone (80 Ci/mmol, New England Nuclear, Boston, MA) dissolved 15 in 100 microliters of "gelatin/phosphate" buffer. buffer contained 0.01 M potassium phosphate (pH 7), 0.82% NaCl, 0.1% NaN3, and 0.2% Difco brand gelatin. testosterone serum sufficient to bind 0.06 pmol 20 testosterone was added in 50 microliters gelatin/phosphate buffer and incubated 1 hour at 370 C. Following an additional 15 minutes in an ice slurry, 400 microliters of dextran/charcoal solution [prepared by adding 1.25 g of powdered charcoal (Sigma, St.Louis, Mo.) 25 to 0.125 g of dextran T-70 (Sigma) and 400 ml of gelatin phosphate buffer] were added to separate the bound and free testosterone. After 15 minutes at 4° C., mixture was centrifuged at 800 x g for 10 minutes and 400 microliters of the supernatant were added to 4 ml of scintillation fluid (Liquiscent, National Diagnostics). 30 The assay measured 0.02-10 pmol of nonlabeled testosterone.

Granulosa cells were isolated from ovaries of rats (21-27 days old) which had been implanted with diethylstilbestrol(DES) "pellets" made by packing solid DES into 1 cm long sialastic tubings. Estrogen treatment has been shown to induce granulosa cell hyperplasia (60) without altering the number of FSH receptors per cell

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Granulosa cells were cultured using a minor the procedure described by of (59) modification Cells were suspended in McCoys Ericksonand Hsueh (62). (GIBCO, Grand Island, NY) and either medium 5a 40,000-50,000 cells 100 microliters containing one milliliter containing 400,000-800,000 cells was added to each well of a standard 96- or 24-well culture plate. The medium also contained penicillin, streptomycin, and Since granulosa cells lack the 100 nM androstenedione. ability to convert C-21 steroids to C-19 steroids (62), the androstenedione was needed as a substrate In addition, androstenedione serves estradiol synthesis. stimulus for progesterone formation (63, although it is not needed for this function. The medium was not changed during the culture interval. Estradiol and progesterone were measured directly from 0.02-0.2 ml culture fluid by RIA using highly specific antisera (65).

dose response curve of LH FIGURE 11 is а receptor mediated steroidogenic activity of hCG/hFSH chimeric beta-subunits co-expressed in COS7 cells with human alpha-subunit, compared with hCG and hFSH. Data from each analog are marked using the nomenclature from TABLE II. Data from analogs with hCG-like activity are plotted with open boxes, rhCG (---), B15 (---), B27 (····); data for rhFSH (---) are plotted with filled boxes; data from analogs with slightly diminished hCGlike activity are plotted with open triangles, B9 (--) and B16 (---); and data from analogs with moderate hCGlike activity are plotted with filled triangles, (--), B17 (---) and B18 (---). The steroidogenic potency of the analogs in this assay is strongly related to their ability their receptor binding activity (see FIGURE 9).

receptor mediated steroidogenic activity of hCG/hFSH chimeric beta-subunits co-expressed in COS7 cells with human alpha-subunit, compared with hCG and hFSH. Data from each analog are marked using the nomenclature from

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TABLE II. Data from analogs with hFSH-like activity are plotted with filled triangles, B11 (--), B17 (---) and B18 (---); data from hFSH (---) are plotted with filled boxes; and data from rhCG (----) are plotted with open boxes. The steroidogenic potency of these analogs in this assay is strongly related to their FSH receptor binding activity (see FIGURE 10).

is known to stimulate adenyl cyclase hCG 10 activity intesticular interstitial (Leydig) cells. accumulates which in response to stimulation of Leydig cells can be used as a measure of the efficacy of hCG or analogs which interact with LH receptors (20). Leydig cells were prepared by digesting 15 rat testes with collagenase as described above. 500,000 cells in a volume of 50 microliters were added to assay tubes containing hormone standard or analog to give a final volume of 100 microliters. This mixture was then incubated at 37° C. for 45 minutes, after which the 20 incubation tubes were placed in a boiling waterbath for 5 minutes and then set on ice. Aliquots of the medium or cAMP standards were then incubated with 3H-cAMP and cyclic AMP dependent protein kinase (Sigma, St. Louis) in an ice slurry overnight. Six hundred microliters of 25 dextran-coated charcoal were added to the samples and incubated 10-15 minutes in an ice slurry. The tubes were centrifuged at 2000 rpm for 15 minutes, 600 microliters of the supernatant were added to 3 ml of scintillation fluid.

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FIGURE 14 is a dose response curve of LH receptor mediated cAMP accumulation in response to hCG and in response to F8 which had been co-expressed in C127 cells with alpha-subunit. This FIGURE also indicates the results of incubating hCG and the F8 analog together. FIGURE 14 illustrates that F8 has a reduced efficacy relative to hCG. In addition, F8 inhibits the ability of hCG to stimulate hCG-induced cAMP accumulation.

All references cited hereinabove and hereinafter, are incorporated by reference.

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The invention being thus described, it will be obvious that the same may be varied in many ways. Such variations are not to be regarded as a departure from the spirit and scope of the invention and all such modifications are intended to be included within the scope of the following claims.

TABLE 1

Examples of Structures of Native Hormones B-subunits

	5	2	30	07	20	8	02	8	8	100	110	120	130	140
55	SKEPLRPRCRPINATLAVEKEGCPVCITVNTTICAGYCPTNTRV	TLAVEKEGCPV	CITVNTTIC	AGYCPINIRV	LOGVLPALPOV	VCNYRDVRFE	SIRLPGCPRG	INPVVSYAVALS	SCOCALCRR	STIDCGGPKD	IPL TCDDPRF	ODSSSSKAPP	PSLPSPSRL	LQGVLPALPQVVCNYRDVRFESIRLPGCPRGVNPVVSYAVALSCQCALCRRSTTDCGGPKDHPLTCDDPRFQDSSSSKAPPPSLPSPSRLPGPSDTP1LPQ
FT.	-W			¥	Ap			.DFP	GP	·	ō-x	rsglifi		
E E	-RGL-QAFT-SS-K	YY	f1-s	S-K	-PVIPMR	1-HELA	-Ah-	-PVIPMRT-HELA-VPD-MFPH-GPL- SRIQAH-PLP-1LFL	149-H-	ex	10AH-P	LP-ILFL		
E.	-RGL	YY	FT-S	s-N-S	HPAAIP	TELA	4	.D-MFP	0 -49-H-	IKVFR	-aA-AP9A	SSS-KDPPSQ	I-LTST-TPT	WPAAIPTELA
<b>E</b>	LGGGGPV-VDQ-MA-TA-GR-REP-	۰۸۵۸-	-MA-TA-	GR-REP-	YRSP-GPPSA-T-GAL-Y-RWA-WI-SD-R-LLPRR- PWA-STVQGLG-AF-GA-GGFGGE	A-T-GAL-Y-	RWA-WI-:	30-R-LLP	RR- P	MA-STVQG	LG-AF-GA-G	GFGGE		
s67.1	GTEC-YGLN-M-11R-D-HGS 1-TL-E-TDL1	-11R-D-HG	S 1-T-	L-E-TOLN	YESTULPRS-GFKEUSY-KVY-ESE- FFIPKD-IK- KTDNDRISMATPS-IVNPLEM	FKEWSY-	KVY-ES-	E- FFIPK	D-IK- K	TONDRIS	MATPS-IVNP	ren		
1113s		S-MQP-QQ-VSLT-LVIQ-PS-H-V-KEP-	-LV19-P	S-H-V-KEP-	FKSPFSTVI-HTY-TDPW-D-H-T-PD-S NMD-STIESLQ-DF-11Q-VLTDGDMW	١١٠	MdQ1	-D-H-T-P	NS-Q	MD-STIES	La-DF-119-	VLTDGDMW		•
hFSH		NS-ELT-1-1-1E-RFS1WY-RDL-	-NIS	Y-RDL-	YKDPARPKI-K	T-TFKELW-	TV-VAHH	YKDPARPKI-KT-TFKELVV-TV-VAHHADSLYT-PTQ-H-GK- DSDSTVRGLG-SY-SFGEMKE	9-H-GK- D	SDSTVRG	LG-SY-SFGE	MKE		-60-
htsh		F-1-TEY-MH1-RRE-AY-L-1M-RDIN	1-1-	M-RD ING	NGKLFLPKYS-DTFIYRTVEILH-A-YFPK-GK- NTDYSIHEAIKTNY-TK-QKSYLVGFSV	TFIYR	TVEILH	.A-YFP	K-GK- N	TDYSIHEA	IKTNY-TK-Q	KSYLVGFSV		-

## elpha-subunits

	10	02	30	0,4	20	3	2	09 . 02	06	
human:	APD VQDCPECTLQENP FFSQPGAP I LQCMGCCFSRAYP I PLRSKK TML VQKNVTSEST CCVAKSYMRV TVMGGFKVENHTACHCST CYYHKS	FFSQPGAP	ILACMGCCFS	RAYPTPLRSK	KTHLVQKNVT	SESTCCVAKS	YMRVTVMGGF	KVENHTACHC	STCYYHKS	
bovine:	FPDGEFTM-GK-KYK-DYEAAFTKANVRE	rK-D		V	-Id	V	FTKAIN	RE	:	
equine:	FPDGEFITK-RKYKL-VY	rKL-V-	λ-		-Id	<b>V</b>	FI	-L1-	IH	
selmon:	YPNSDKTNMGKP-T1-PN -M-1A	Nd-I	-HTH-	····¥·····	1p1	3¥	GE170	p-TE	:	

those in the respective hCG subunits are indicated with hyphens (-). The gaps present in the sequences in the figure are due to differences in the number of emino acids in intracysteine regions between TSH and the other 8-subunits, and salmon alpha-subunit and the other alpha-subunits. Numbers refer to position Amino acid sequences of glycoprotein hormone subunits, hCGB (40), hLHB (80), bLHB (81,83), eLHB (79), cLHB (78), sGTHIB and sGTHIIB (70), hFSHB (77), hTSHB. (82), human-alpha (43), bovine-alpha (84), equine-alpha (85) and salmon-alpha (70). Single letter amino acid code is used (Afalanine, Cacysteine, Daspartic acid, Esglutamic acid, Fephenylalanine, Geglycine, Hehistidine, Irisoleucine, Kelysine, Leleucine, Memethionine, Neasparagine, Peproline, Geglutamine, Ranginine, Saserine, Tathreonine, Vavaline, Watryptophan, Yatyrosine). The B-subunits are from human (h), bovine (b), equine (e), chicken (c) and salmon (s). The sequences are aligned to facilitate examination of homology with either hCG B-subunit (top) or hCG alpha-subunit (bottom). Reidues identical to in hGG alpha-subunit and B-subunit sequences, other hormone subunits are aligned by their cysteine residues. TABLE 1

Examples of hCG/hFSH CHIMERAS

			11-KT-TFKELV	L-YKOPARPK1-KT-TFKELVY-TV-VAHHADSLYT-PTG-H-GK-DSDSTVRGLG-SY-SFGEMKE	HADSLYT-P-	-T9-H-GK-D	NISOS	RGLG-SY-SF	GENKE		
=	NS-ELT-1-1-1										į
85	W									,	:
83	Y-RD	-RDL									:
96		YKDPARPKI-					:				:
82			KT-T				:				:
98	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0		FKELVY-TV-V-	r-TV-V							:
87				HVAH	AHHADSLYT-P						:
88						10-H-GK					:
8						Q	soso				i
B10							TV	TVRGLG-SY-SFGEMKE	GEMKE		-
118						Q	DSDS TV	TVRGLG-SY-SFGEMKE	GENKE		61
812							VI	TVRGLG-SY-SFGENKE	GENKE		- :
813						0	V1S0S0	TVRGLG-SY-SFGEMKE	GEMKE		:
814			KT-T								;
815		Y-RDL-YKDPARPKI-KT-T-	(I-KT-T								:
816			(I-KT-I			0	soso				:
817	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	Y-RDL-YKDPARPKI-KT-T	(1-KT-T			<b>0</b>		TVRGLG-SY-SFGEMKE	GEMKE		
818						0	VISOSO	TVRGLG-SY-SFGE	w		
919		-RDL-YKDPARPKI-KT-T	(I-KT-T			0	DSDSTV	TVRGLG-SY-SFGQ	3		
820							1	TVRGLG-SY			:
<b>B21</b>							1	TVRSFGE	<b>8</b>		
822	NS-ELT-1-1-1E-RFSIWY-RD	-ROL-YKOPARPI	IL-YKDPARPKI-KT-TFKELVY-TV-VAHHADSLYT-PTQ-H-GK	1-TV-VAH	HADSLYT-P-	-10-H-GK	:	SY-SFGEMKE	GEMKE		
823							•	GLG-SY-SFGE	띯		
<b>B</b> 24	Y-RDL-YKDPARPKI	-RDL-YKOPARPI	D			•					:
825	NS-ELT-1-1-1E-RFS1WY-RD	-RDL-YKOPARPI	IL-YKDPARPKI-KT-TFKELVY-TV-VAHHADSLYT-PTQ-H-GK-DSDS-	Y-TV-VAH	HADSLYT-P-	TQ-H-GK-D	sosc				:
<b>B</b> 26	NS-ELT-1-1-1-1E-RFSIWY-RDL-YKDPARPKI-KT-1FKELVY-TV-VAHHADSLYT-PTQ-H-GK-DSDS-	-RDL-YKDPARPI	KI-KT-TFKELM	Y-TV-VAH	HADSLYT-P-	TQ-H-GK-D	SOSC	SY-SFGENKE	GENKE		

TABLE 111

# Examples of hCG/hTSH CHIMERAS

		9	20	ន្ត	9	20	<b>%</b>	2	88	8	<b>100</b>	110	120	130	140
hcgs		SKEPLRPRCRP1NATLAVEKEGCPVC11VNTT1CAGYCPTM1	VEKEGCPVC	ITVNTTICAG	YCPIMTRVLQ	GVLPALPQVV	CNYRDVRFES	IRLPGCPRGV	HPVVSYAVAL!	COCALCRR	RVLQGVLPALPQVVCNYRDVRFESIRLPGCPRGVNPVVSYAVALSCQCALCRRSTTDCGGPKDHPLTCDDPRFQDSSSSKAPPPSLPSPSRLPGPSDTP1LPQ	PLTCDOPRF	XOSSSSKAPPF	SLPSPSRLP	GPSDTP1LPQ
hTSHB	•	F-I-TEY-MHI-RRE-AY-L-IM-RD	-RRE-AY-		M-RD INGK	XPKYS-D-	-TFIYRT	VE1LH-	A-YFP	-K-GK- N	)INGKXPKYS-DTFIYRTVEILH-A-YFPK-GK- NTDYSIHEAIKTNY-TK-GKSYLVGFSV	KTNY-TK-9	(SYLVGFSV		
5		F-1-1EY-MHI													
8			RRE-AY-	١٠٠١	•										
ន		18-18			=	XPKYS-D-	-11-				NGKXPKY-S-DT				
ಕ				1		•	FIYRT	VE1							
ស		0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0			1 1 1 1 1 1			¥7	A-YFP		LH-A-YFP				-
8			1							N -K-GK-	K-GK- NIDVS				62-
<b>C</b> 2			4									KTNY			
5											TK-TK-TK-TK-TK-TK-TK-TK-TK-TK-TK-TK-TK-T	D-X1	(SYLVGFSV		
ಬ			0 0 0 0 0							-K-GK- N		KTNY-TK-Q	(SYLVGFSV		

See Table I for legend. In this particular table, "X" denotes the sequence LFL from hISHB.

TABLE 1V
Examples of hcg/blH CHIMERAS

		0	20	R	07	20	3	20	8	8	100	110	120	130	140
hcg8	SKEPLRPR(	CRPINATL	.AVEKEGCPV	SKEPLRPRCRP I NATLAVEKEGCPVC I TVNTT I CAGYCP TI	GYCPTMTRVLQ	MTRVLQGVLPALPQVVCHYRDVRFESIRLPGCPRGVHPVVSYAVALSCQCALCRRSTTDCGGPKDHPLTCDDPRFGDSSSSKAPPPSLPSPSRLPGPSDTP1LPQ	CNYRDVRFE	SIRLPGCPRG	INPVVSYAVAL	SCOCALCRRS	ТТОСССЕРКОН	PL.TCDOPRF9	DSSSSKAPPP	SLPSPSRLP	SPSDTP1LPQ
PL#8			AA-	FT-S	-RGL-QAFT-SS-KPVIPMRT-HELA-VH-GPL-SRTQAH-PLP-1LFL	VIPMR-	-T-HELA	dh-	.D-MFP	#-GPL-	SRT	0AH-PL	P-11.FL		
2	-RGL·	ō-	YY-	-RGL-QAA											
62			YY-		:						i				
20						VIPMR-								•	
70						1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	T-HELA-V	····							
92								4	PD-MFP						
90											-S-				-6
20						1			·S. ]	· · · · · · · · · · · · · · · · · · ·	÷				53-
80											•	RTQAH			
6	-RGL		YY	FT-S	-RGL-QAA	VIPMR-				: : : :	i		P-ILFL		
010	-RGL		YY	FT-S	-RGL-QAAFT-SS-KPVIPMRT-HELA-VPD-MFPH-GPL-SRTQAH-PLP-ILFL	VIPMR-	-T-KELA	-dA-	.D-IKFP	H-GPL·	.SRT	QAH-PL	P-11FL		
110									•	-169-H	H-GPL-SRTQAH-PLP-1LFL	0AH-PL	P-1LFL		
012							•	1		• • • • •	Ex	RTQAH-PLP-1LFL	P-11FL		

See Table 1 for legend.

TABLE Y EXAMPLES OF HCG/CLH CHIMERAS

	10	20	R	07	20	8	20	80	8	100	110	120	130	140	
hcas	hcgb skeplrprcrpinatlavekegcpucituntticagycptm	INATLAVEKI	EGCPVC1 TVNT	TICAGYCPTHT	TRVLQGVLPALP	QVVCNYRDVR	FESTRLPGCPI	RGVMPVVSYA	/ALSCOCALC	RRSTTOCGGP	OHPL TCOOP	RFODSSSSKA	PPPSLPSPSR	ITRVL GGVLPAL PQVVCNYRDVRFES IRL PGCPRGVNPVVSYAVAL SCQCAL CRRST TDCGGPKDHPL TCDDPRFQDSSSSKAPPPSL PSPSRL PGPSDTP I LPQ	
CLHB	-RGL	V	-A1FT-	A-SS	/MPAAI	bTEL	V-	PD-HFP		01KV	FR-9A-AP	aasss-kopp	.SQ-LTST-TP	elhb -RGLAIFT-SS-VMPAAIPTELA	_
<b>=</b>			0 6 8 8 8	N-S	/HPAAI					:					
23										91K					
<b>5</b>										۸	FR-QA			VFR-QA	
73	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1					7 8 4 9 9					dV	Y0	1 1 1 1 1 2	APQA	
æ									1	91KV	FR-QA-AP	V			
	•	-												(	-6
998	See Table 1 for legend.	egend.												) •	

IABLE VI

Examples of hCG/hLH CHIMERAS

	10	50	8	07	22	3	2	88	8	100	110	120	130	140
hcgs	SKEPLRPRCRP I NATLAVEKEGCPVCI TVNTTI CAGYCPT	ATLAVEKEGCP\	CITVNTTIC	<b>LGYCPTMTRVL</b> (	<b>JGVLPALPQV</b>	ICHYRDVRFES	IRLPGCPRG	INPVVSYAVAL	SCACALCRRS	TTDCGGPKDH	PLTCOOPRFQ	MTRVLGGVLPALPQVVCNYRDVRFESIRLPGCPRGVNPVVSYAVALSCQCALCRRSTTDCGGPKDHPLTCDDPRFQDSSSSKAPPPSLPSPSRLPGPSDTP1LPQ	SLPSPSRLP	PSDTP1LPQ
ht.HB	-RW-HI			<b>I</b>	-Ap			.DFP	R-GP		10-H	SGLLFL		
۲					1	1			- ;					
<b>4</b> 5				•			0 0 0 0 0 0		1					
S.								DFP S	R-GP					
A38					3 4 3 6 6 4									
A3b				1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		1		8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	R-GP					
<b>y</b> ¢								1 1 1 0 0 0 0 0			SB-GLLFL	SGLLFL		-
AS				1	dY									65-
<b>Y</b> ę	N-N			•				1 2 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0						
<b>V</b>				<b>=</b>	dH			DFPR-GP S	R-GP					
<b>A8</b>							1	BFPR-GP		S		H-OLSGLLFL		
8						1		R-GPR-GPR-GPR-GPR-GP	R-GP	S				
A10				¥					R-GP S	S			1	

See Table I for legend.

See Table I for legend. Deletions are denoted by empty spaces in place of hGGs residues.

TABLE VIII

Examples of Other Analogs

SKEPLRPRCRPINATLAVEKEGCPVCITVNTTICAGYCPT0 -M
T T

See Table I for legend. In G9, "X" = any residue except Pro, "J" = Ser or Thr

TABLE

Examples of alpha-subunit chimeras and mutants

APDVQ	10 20 30 40 50 60 70 80 90 90 APDVQDCPECTLQENPFFSQPGAPILQCMGCFSRAYPTPLRSKKTMLVQKNVTSESTCCVAKSYNRVTVMGGFKVENHTACHCSTCYYHKS	20 FFSQPGAPILO	30 ICHGCCFSRA	40 YPTPLRSKKT	50 HLVQKNVTSE	60 Stccvaksyn	70 RVIVMGGFKVE	80 ENHTACHCST	90 CYYHKS
FPDGEFIM.GK-KKYK-DY	(-KK	/K-Dγ-		Y	b1	AAFT	KANVR	3	:
¥	-KK	·····K·K··KY··K·D···Y····		<b>y</b>	[d	AAFT	A	ü	;
K-KKYK-DY	-KK	rK-0Y-							:
				¥	Jd	AAFT	APIAAFTKANVR	-	:
FPDGEFTM-G									:
FPDGEF1M-GAFTKANVRE				γ	ld	AAFT	KANVR	ų.	:
FPDGEFTM-GK-KKYK-DY	.KK	/K-DY-							:
FPDGEFIM-GK-DYAPIAAFTKANVR		···K-DY-		¥	<b>b l</b>	AAFT	KANVR	-	:
FPDGEFTM-GK-KKYK-DYA	ή. -Κ	/K-DY-			Id	V	pIAEE		:
FPDGEFTM-GK-KKY	.κ κ		1	¥	b1	AAFT	AEEE		:
FPDGEFTM-GK-KKYK-DY	.κκ	/K-DY-		Y	P1AAFTKA	AAFT	X	<b>4</b>	:
FPDGEFTM-G		κ.Dγκ					* * * * * * * * * * * * * * * * * * *		:
FPDGEFTM-G		X-D		V			NVR	Ü	:
FPDGEFTM-G				¥	VJd	· · · · · · · · · · · · · · · · · · ·	KVR	•	
						: : : : : : : :			_
				1				¥	:
									H1
									HHH
			#S#	AYPTPLRSKI	KTMLVQKNVT	SESTCCVAKS	RSRAYPTPLRSKKTMLV9KNVTSESTCCVAKSYNRVTVMGGFKVENHTACHCSTCYYHKS	VENHTACHOS	STCYYHKS
FPDGEFTM-G	-					AFT	AFTKARVRE-	-	:

See Table 1 for legend. The gap in H19 is a deletion mutation.

TABLE X

**Examples of Fusion Constructs** 

	g	ohe)	~			<b>ب</b>		-69 <b>-</b>
140	TRVLQGVLPALPQVVCNYRDVRFESIRLPGCPRGVNPVVSYAVALSCQCALCRRSTTDCGGPKOHPLTCODPRFQDSSSSKAPPPSLPSPSRLPGPSDTPILPQ	(hCG-alpha)(hCG-alpha)	(NSN-G)	(1961)	,(CAT) <sup>4</sup>	Strep) <sup>5</sup>		
136	(PPPSLPSPSR			(196)	(CA1) <sup>4</sup>		tein]	
120	PRFODSSSKU					1	second pro	
110	PKOHPLTCODE						[second protein]	
100	CRRSTTDCGG				•		:	
8	AVAL SCOCAL							
08	PRGVNPVVSY							
02	RFESIRLPGC						•	•
3	POVVCNYRDV							
20	RVLQGVLPAL							
07	ICAGYCP THT							
93	CPVC! TVN!T							
50	INATLAVEKEG							
10	SKEPLRPRCRPINATLAVEKEGCPVCITVN1T1CAGYCPTM1						•	(X) [Z]
	PCG8	5	75	2	75	35	<b>J</b>	75

See Table I for legend.

Fusion protein with entire hCG alpha-subunit amino acid residues -24 to +92 (see Figure 2).

Fusion protein with transmembrane and cytoplasmic domains of vesicular stomatitis virus (VSV) G- protein.

Fusion protein with hoos in place of immunoglobulin variable region.

Fusion protein with chloremphemical acetyltransferase.

5 Fusion protein with streptavidivin.

6 Fusion protein with hoos residues 9-110 plus any of the proteins in J1-5.

7 Fusion protein with any glycoprotein hormone 8-subunit [X] plus a second protein [2] such as an enzyme or binding protein.

TABLE XI
MONOCLONAL ANTIBODY BINDING TO HUMAN/BOVINE atpha-SUBUNIT

CHIMERAS CO-EXPRESSED WITH hCG B-SUBUNIT

	body	A112 A202 A501	++ ++	**			. ++	+	*	+	- ++	•		nt ++ ++	nt ++ +	**
	Monoclonal Antibody	A109	‡	‡		•	•	٠	*	+	٠	•		‡	+	‡
	Monoclo	A102	:	*		•		<b>‡</b>	*	*	•	‡	•	‡	*	‡
		A101	‡	*	•			*	*	‡		+		Ę	겉	;
nit		<b>B</b> 109	:	:	*	<b>‡</b>	:	*	:	*	:	:	:	:	*	:
alpha-subunit	or Analog		ohce	rhCG	b/+hcg8	£	H2	13	7.1	#5	H6	H7	£	111	H12	H14

Data are presented as binding of labeled antibody to chimeric proteins relative to binding to urinary hGG; ++++  $\pm$  greater than 200% cpm bound by hCG standard, +++  $\pm$  greater than 110%, ++  $\pm$  50%-110%, +  $\pm$  10%-69%, -  $\pm$  less than 10% and nt  $\pm$  not tested.

ngaman aman colombana mara managa colombana (marama).

TABLE XII

Monoclonal Antibody Binding to Membranes from COS7 Cells Expressing hCGB/VSV-G Fusion Proteins +/- /-Subunit

	Transfection	ction	Ratio	Ratio MAb/MAb+hCG	o+hcG	·
elphe	alpha+8"tail"1	B"tail"	МАБ	alpha+B <sup>1</sup>	-#a	B1
A111	9269	952				
A111 + 1 /g hCG	1024	1178	A111	A111 6.8 0.8	0.8	
B109	9509	1998				
B109 + 1 /g hcG 1344	1344	1514	8109	8109 4.5 1.3	1.3	
8105	11872	26104				
8105 + 1 /g hcc 1049	1049	1268	8105	8105 11.3 20.6	20.6	

1. B analog J2 (hCGB/VSV-G fusion protein)

TABLE XIII MONOCLOMAL ANTIBODY BINDING TO CHIMERIC hcg/hfsh

B-SUBUNITS CO-EXPRESSED WITH ALpha-SUBUNIT

	A109	‡	+	‡	nt	•	•	•	at	*	•	nt	돧
ntibody	1098	•	nt	6	•	at T	‡	‡	*	•	*	•	*
Monoclonal Antibody	B109	<b>‡</b>	‡	•	٦ţ	•	•	•	at	•	•	aţ	
Morro	1018	<b>‡</b>	aţ	*	<b>‡</b>	Ę	•	•		<b>‡</b>	•	<b>‡</b>	٠
	A113	<b>‡</b>	‡	:	:	<b>‡</b>	*	:	:	:	:	*	:
B-Subunit	or Analog	rhcG	83	68	118	914	815	816	917	818	828	827	rhFSH

Data are presented as antibody binding to chimeric proteins relative to binding to hCG or hFSH standard; ++ = 50x-110x, + = 10x-50x, - = less than 10x, nt = not tested

COMPOSITE CMES

TABLE XIV BINDING OF HUMAN/BOVINE alpha-Subunit Chimeras Co-expressed

WITH hCG B-SUBUNIT TO LH RECEPTORS

BioIRMA	B105	‡		‡	‡	‡	‡	<b>‡</b>	‡	‡	*	<b>‡</b>
alpha-Subunit	or Analog	פטעה	rhcG	85/+hcG8	1.1	. 24	H3	7±	H5	H6	ZH	. H.

Data are presented as binding of mAb B105 to hormone analog/receptor complex relative to binding to hCG/receoptor comlex; ++ = 50X-150X, + = 10X-49X, = less

than 10%

TABLE XV

MONOCLONAL ANTIBODY BINDING TO STABLY TRANSFECTED COS7 CELLS

EXPRESSING hCG B-SUBUNIT OR hCGB/VSV-G

Cell	8105	8105+300ng hcg	Ratio -/+ hCG
cos7-hcg8	4772	5544	2.1
COS7-hcG8/vsv-G	45762	1887	24.3
(75)			

-75-

We claim:

1. A physiologically active glycoprotein hormone analog which has coupled <u>alpha</u> and <u>beta</u>-subunits which glycoprotein has bonding affinity to two receptors of different native glycoprotein hormones, with the proviso that the analog is not human LH-<u>beta-CG-beta</u> wherein the <u>beta-subunit</u> binds to a common gonadal receptor.

10

5

2. A physiologically active glycoprotein hormone analog which has coupled <u>alpha</u> and <u>beta-subunits</u> which has binding affinity to two different receptors.

15

3. A physiologically active glycoprotein hormone analog which has coupled <u>alpha</u> and <u>beta-subunits</u> which has binding affinity to two receptors of different native glycoprotein hormones with the proviso that either one of the subunits is a species other than human.

20

4. The physiologically active glycoprotein hormone analog according to claim 2, wherein the <u>beta-subunit</u> contains amino acid sequences found in native CG and FSH.

25

5. The physiologically active glycoprotein hormone analog according to claim 2, wherein the <u>beta-</u>subunit contains amino acid sequences found in native CG and TSH.

30

6. The physiologically active glycoprotein hormone analog according to claim 4, wherein at least one of the subunits is human.

35

7. The physiologically active glycoprotein hormone analog according to claim 5, wherein at least one of the subunits is human.

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8. The physiologically active glycoprotein hormone analog according to claim 6, wherein the <u>beta-</u>subunit is human.

- 9. The physiologically active glycoprotein hormone analog according to claim 7, wherein the <u>beta-</u>subunit is human.
- 10. The physiologically active glycoprotein hormone analog according to claim 3, wherein the subunit is other than human and is selected from the group consisting of bovine, equine, ovine, and pisces.
- 11. The physiologically active glycoprotein hormone analog according to claim 10, wherein the <u>beta-</u>subunit is other than human.

# 1/16 Figure 1. hCG 8-subunit cDNA and protein sequence

1		10		2	0			30			40			!	50			60		
5'-AGA	CAAG	GCAG	GGGA	CGCA	CCAA	GG	ATG	GAG	ATG	TTC	CAG	GGG	CTG	CTG	CTG	TTG	CTG	CTG	CGT	AGC-3
3'-TCT	GTTC	CGTC	CCCT	GCGT	GGTT	CC	TAC	стс	TAC	AAG	GTC	ССС	GAC	GAC	GAC	AAC	GAC	GAC	GAC	TCG-5'
							H	ε	M	F	Q	G	L	L	L	L	L	L	L	s
						•	-20										-10			
													٠							
70			į	80			90			100			11	10			120			130
5'-ATG	GGC	GGG	ACA	TGG	GCA	TCC	AAG	GAG	CCG	CTT	CGG	CCA	CGG	TGC	CGC	CCC	ATC	AÄT	GCC	ACC-3
3'-TAC	CCG	CCC	TGT	ACC	CGT	AGG	TTC	стс	GGC	GAA	GCC	GGT	GCC	ACG	GCG	GGG	TAG	TTA	CGG	TGG-5'
	G	G	Ţ	u	A	s	ĸ	ε	Ρ	L	R	P	R	С	R	P	I	N	A	T
					-1	1									10					
													Hine	:11					Nac	<u>. I.</u>
		14	40			150			160			17		11_	1	180			<u>Nac</u> 190	<u> </u>
5'-CTG	GCT		-	AAG		-	TGC	ccc		TGC	ATC		70				ATC	TGT	190	GGC-3'
		GTG	GAG		GAG	GGC			GTG			ACC	TO GTC	AAC	ACC	ACC	TAG	ACA	190 GCC	GGC-3'
		GTG	GAG	TTC K	GAG	GGC			GTG			ACC	TO GTC	AAC TTG	ACC	ACC			190 GCC	GGC-3'
3'-GAC		GTG	GAG CTC	TTC	GAG CTC	GGC	ACG		GTG	ACG	TAG	ACC TGG	O GTC CAG	AAC TTG	ACC TGG	ACC TGG	TAG	ACA	190 GCC	GGC-3'
3'-GAC		GTG	GAG CTC	TTC K	GAG CTC	GGC	ACG	<b>GGG</b> P	GTG	ACG	TAG I	ACC TGG T	O GTC CAG	AAC TTG	ACC TGG	ACC TGG T	TAG	ACA	190 GCC	GGC-3'
3'-GAC	CGA A	GTG	GAG CTC	TTC K 20	GAG CTC	GGC	ACG	<b>GGG</b> P	GTG CAC V	ACG C	TAG I	ACC TGG T	GTC CAG V	AAC TTG N 30	ACC TGG	ACC TGG T	TAG	ACA	190 GCC	GGC-3'
3'-GAC L	CGA A	GTG CAC V	GAG CTC E	TTC K 20	GAG CTC E	GGC CCG G	C	PS1	GTG CAC V	ACG C	TAG I MI 23	ACC TGG T	GTC CAG V	AAC TTG N 30	ACC TGG T	ACC TGG T	TAG I 1	ACA C	190 GCC CGG A	GGC-3'
3'-GAC	CGA A 20 TGC	GTG CAC V	GAG CTC E	TTC K 20	GAG CTC E 210 ACC	GGC CCG G	ACG C	PS1 220 CTG	GTG CAC V	ACG C Ppc	TAG  I  AI  GTC	ACC TGG T Na	TO GTC CAG V	AAC TTG N 30	ACC TGG T	ACC TGG T  Bsi	TAG I 2361 CAG	ACA C 250 GTG	190 GCC CGG A	GGC-3' CCG-5'
3'-GAC	CGA A 20 TGC	GTG CAC V	GAG CTC E	TTC K 20	GAG CTC E 210 ACC	GGC CCG G	ACG C	PS1 220 CTG	GTG CAC V	ACG C Ppc	TAG  I  AI  GTC	ACC TGG T Na	TO GTC CAG V	AAC TTG N 30	ACC TGG T	ACC TGG T  Bsi	TAG I 2361 CAG	ACA C 250 GTG	190 GCC CGG A	GGC-3' CCG-5' G
3'-GAC L 5'-TAC 3'-ATG	CGA A 20 TGC ACG	GTG CAC V	GAG CTC E	TTC K 20	GAG CTC E 210 ACC TGG	GGC G CGC GCG	ACG C	PS1 220 CTG	GTG CAC V CAG	ACG C Ppc GGG CCC	TAG  I  AI  GTC	ACC TGG T Na	TO GTC CAG V	AAC TTG N 30	ACC TGG T	ACC TGG T  Bsi	TAG I 2361 CAG	ACA C 250 GTG	190 GCC CGG A	GGC-3' CCG-5' G TGC-3'

## 2/16 Figure 1. hCG B-subunit cDNA and protein sequence (continued)

TaqI 270 280 290 300 310 260 5'-AAC TAC CGC GAT GTG CGC TTC GAG TCC ATC CGG CTC CCT GGC TGC CCG CGC GGC GTG AAC CCC-3' 3'-TTG ATG GCG CTA CAC GCG AAG CTC AGG TAG GCC GAG GGA CCG ACG GGC GCG CAC TTG GGG-5' G C P RFESI 60 Pvu II Apa LI 340 350 360 370 320 5'-GTG GTC TCC TAC GCC GTG GCT CTC AGC TGT CAA TGT GCA CTC TGC CGC CGC AGC ACC ACT GAC-3' 3'-CAC CAG AGG ATG CGG CAC CGA GAG TCG ACA GTT ACA CGT GAG ACG GCG GCG TCG TGG TGA CTG-5' YAVALSCQCA CRRSTTD 90 80 PpuMI 400 410 420 430 440 390 5'-TGC GGG GGT CCC AAG GAC CAC CCC TTG ACC TGT GAT GAC CCC CGC TTC CAG GAC TCC TCT TCC-3' 3'-ACG CCC CCA GGG TTC CTG GTG GGG AAC TGG ACA CTA CTG GGG GCG AAG GTC CTG AGG AGA AGG-5' C 110 120 100 Apai Sma I 490 450 460 470 480 500 5'-TCA AAG GCC CCT CCC CCC AGC CTT CCA AGC CCA TCC CGA CTC CCG GGG CCC TCG GAC ACC CCG-3' 3'-AGT TTC CGG GGA GGG GGG TCG GAA GGT TCG GGT AGG GCT GAG GGC CCC GGG AGC CTC TGG GGC-5' L P G P S T 130 140 530 539 510 520

5'-ATC CTC CCA CAA TAA AGGCTTCTCAATCCGC(A)40-3'
3'-TAG GAG GGT GTT ATT TCCGAAGAGTTAGGCG(T)40-5'

.\_\_\_\_.

## 3/16 Figure 2. hCG alpha-subunit cDNA and protein sequence

Ncol Hincll 5'-CAGTCAACCGCCCTGAACACATCCTGCAAAAAGCCCAGAGAAAGGAGCGCC ATG GAT TAC TAC AGA AAA TAT-3' 3'-GTCAGTTGGCGGGACTTGTGTAGGACGTTTTTCGGGTCTCTTTCCTCGCGG TAC CTA ATG ATG TCT TTT ATA-5' M D Y Y R K Y -24 -20 100 5'-GCA GCT ATC TIT CTG GTC ACA TTG TCG GTG TTT CTG CAT GTT CTC CAT TCC GCT CCT GAT-3' 3'-CGT CGA TAG AAA GAC CAG TGT AAC AGC CAC AAA GAC GTA CAA GAG GTA AGG CGA GGA CTA-5' A A I F L V T L S V F L H V L H S A P D - 1 -10 150 BsmI 5'-GTG CAG GAT TGC CCA GAA TGC ACG CTA CAG GAA AAC CCA TTC TTC TCC CAG CCG GGT GCC-3' 3'-CAC GTC CTA ACG GGT CTT ACG TGC GAT GTC CTT TTG GGT AAG AAG AGG GTC GGC CCA CGG-5' V Q D C P E C T L Q E N P F F S Q P G A 10 250 200 Xbal 5'-CCA ATA CTT CAG TGC ATG GGC TGC TGC TTC TCT AGA GCA TAT CCC ACT CCA CTA AGG TCC-3' 3'-GGT TAT GAA GTC ACG TAC CCG ACG ACG AGA GAT TCT CGT ATA GGG TGA GGT GAT TCC AGG-5' PILQCMGCCFSRAYPTPLRS 30 40 300 5'-AAG AAG ACG ATG TTG GTC CAA AAG AAC GTC ACC TCA GAG TCC ACT TGC TGT GTA GCT AAA-3' 3'-TTC TTC TGC TAC AAC CAG GTT TTC TTG CAG TGG AGT CTC AGG TGA ACG ACA CAT CGA TTT-5' K K T M L V Q K N V T S E S T C C V A K 50 350 Dralli PstI 5'-TCA TAT AAC AGG GTC ACA GTA ATG GGG GGT TTC AAA GTG GAG AAC CAC ACG GCG TGC CAC-3' 3'-AGT ATA TTG TCC CAG TGT CAT TAC CCC CCA AAG TTT CAC CTC TTG GTG TGC CGC ACG GTG-5' S Y N R V T V M G G F K V E N H T A C H 70 80 400 Scal 5'-TGC AGT ACT TGT TAT TAT CAC AAA TCT TAA ATGTTTTACCAAGTGCTGTCTTGATGACTGCTGATTTTC-3' 3'-ACG TCA TGA ACA ATA ATA GTG TTT AGA ATT TACAAAATGGTTCACTACAGAACTACTGACGACTAAAAG-5' CSTCYYHKS\*\*\* 90 92 500 5'- TGGAATGGAAAATTAAGTTGTTTATGGCTTTGTGAGATAAAACTCTCCTTTTCCTTACCATACCACTTTGAC-3' 3'- ACCTTACCTTTTAATTCAACAAATCACAAATACCGAAACACTCTATTTTGAGAGGAAAAGGAATGGTATGGTGAAACTG-5' PstI 5'- ACGCTTCAAGGATATACTGCAGCTTTACTGCCTTCCTCCTTATCCTACAGTACAATCAGCAGTCTAGTTCTTTTCATTT-3' 3'- TGCGAAGTTCCTATATGACGTCGAAATGACGGAAGGAGGAGTATAGGATGTCATGTTAGTCGTCAGATCAAGAAAAGTAAA-5' 600 HindIII 5'-GGAATGAATACAGCATTAAGCTT-3' 3'-CCTTACTTATGTCGTAATTCGAA-5'

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# 4/16 FIGURE 3. Oligonucleotides used to construct pKBM and pKBMt

3a pUC18 / pKBMt

HindIII

Xbai

XhoI

5'-AGCTTCTCGAGT-3'

3'-AGAGCTCAGATC-5'

3b pKBMt / pKBM

BamHI

Sali

EcoR1

BasHII SacII

5'-GATCCGCGCGCGTCGACCCGCGGG-3'

3'-GCGCGCGCAGCTGGGCGCCCTTAA-5'

# 5/16 Figure 4a. hCGB' DNA and protein sequence

1 5'-AGA		10		20	3			30			40			9	50			60		
	CAAG		GGGA			GG.	ATG	GAG	ATG	TTC	CAG	GGG	CTG	CTG	CTG	TTG	CTG	CTG	CGT	AGC-3'
3'-TCT													GAC							TCG-5'
J .C.						-	M	Ε	М	F	Q	G	L	L	L	Ł	L	L	L	s
							-20										- 10			
70				30			90			100			1	10		•	120			130
	GGC	GGG			GCA	TCC	AAG	GAG	CCG	CTT	CGG	CCA	CGG	TGC	CGC	CCC	ATC	AAT	GCC	ACC-3'
3/-TAC				ACC									GCC							TGG-5'
H	G	G	T	¥	A	s	K	E	P	L	R	P	R	C	R	₽	1	N	A	T
••	•	-	•		-1	1									10					
					-															
													Hin	11:					Na	<u> </u>
			40			150			160				70			180			190	
5'-CTG	GCT	GTG	GAG	AAG	GAG	GGC	TGC	CCC	GTG	TGC	ATC	ACC	GTC	AAC	ACC	ACC	ATC	TGT	GCC	GGC-3'
3'-GAC	CGA	CAC	СТС	TTC	CTC	CCG	ACG	GGG	CAC	ACG	TAG	TGG	CAG	TTG	TGG	TGG	TAG	ACA	CGG	CCG-5'
L	A	٧	Ε	K	E	G	C	P	V	C	I	Ţ	٧	N	T	T	I	Ć	A	G
				20										30						
								Psi	11		/	/				Bst	<u> 1361</u>	-		
	_	00			210			220				50			240			250		
5'-TAC	TGC																			TGC-3'
3'-ATG	ACG	GGG	TGG	TAC	TGG	GCG	CAC	GAC									GTC			ACG-5'
Y	C	P	T	H	T	R	٧	L	Q	G	٧	L	Р	A	L	P	Q	٧	٧	С
			40										50							
						_														
_						Tac	•			20	•		-	500			310			
_	50			270		***	280	700	476	-	90 CTC	CCT			cce	ccc		GTG	440	ccc-3'
					GCG															GGG-5'
3'-TTG	AIG Y	GCG R	D	V	B B	F	E	S	170	R	L	P	G	C	P	R	G	V	N	P
N	•	60	U	•	•	•	_	3	•	_	-	70	•	Ū	•	•	•	•		•
		-																		
							P۱	vu I	i		Api	a LI								
320		;	330			340	P	vu I		50	Apa		360			370			3	B0
	GTC	_		GCC	GTG				3!		,	3		TGC	CGC		AGC	ACC	_	BO GAC-3'
		TCC	TAC			GCT	стс	AGC	3! TGT	CAA	TGT	GCA	CTC	TGC ACG		CGC		ACC TGG	ACT	
5'-GTG		TCC	TAC			GCT	стс	AGC	3! TGT	CAA	TGT	GCA	CTC			CGC			ACT	GAC-3'
5'-GTG 3'-CAC	CAG	TCC AGG	TAC ATG			GCT	CTC GAG	AGC TCG	3! TGT ACA	CAA GTT	TGT ACA	GCA CGT	CTC GAG	ACG	GCG	CGC	TCG	TGG	ACT TGA	GAC-3' CTG-5'
5'-GTG 3'-CAC	CAG V	TCC AGG	TAC ATG			GCT	CTC GAG	AGC TCG	3! TGT ACA	CAA GTT	TGT ACA C	GCA CGT	CTC GAG	ACG	GCG	CGC	TCG	TGG	ACT TGA	GAC-3' CTG-5'
5'-GTG 3'-CAC	CAG V 80	TCC AGG S	TAC ATG Y		CAC V	GCT CGA A	CTC GAG	AGC TCG S	TGT ACA C	CAA GTT	TGT ACA C 90	GCA CGT A	CTC GAG	ACG	GCG R	CGC GCG R	TCG	TGG	ACT TGA T	GAC-3' CTG-5'
5'-GTG 3'-CAC V	CAG V 80	TCC AGG S Ppul	TAC ATG Y	CGG A	CAC V 400	GCT CGA A	CTC GAG L	AGC TCG S	3! TGT ACA C	CAA GTT Q	TGT ACA C 90	GCA CGT A	CTC GAG L	ACG C	GCG R 430	CGC GCG R	TCG S	TGG T	ACT TGA T	GAC-3' CTG-5' D
5'-GTG 3'-CAC V	CAG V 80	TCC AGG S Ppul 390 GGT	TAC ATG Y	CGG A	CAC V 400 GAC	GCT CGA A	CTC GAG L	AGC TCG S	TGT ACA C	CAA GTT Q	TGT ACA C 90	GCA CGT A 420 GAC	CTC GAG L	ACG C	GCG R 430 TTC	CGC GCG R	TCG S	TGG T 4. TCC	ACT TGA T	GAC-3' CTG-5' D
5'-GTG 3'-CAC V 5'-TGC 3'-ACG	CAG V 80 GGG CCC	TCC AGG S Ppul 390 GGT CCA	TAC ATG Y	AAG TTC	CAC V 400 GAC CTG	GCT CGA A	CTC GAG L CCC GGG	AGC TCG S	TGT ACA C	CAA GTT Q TGT ACA	TGT ACA C 90	GCA CGT A 420 GAC CTG	CTC GAG L CCC GGG	ACG C	GCG R 430 TTC AAG	CGC GCG R CAG GTC	TCG S GAC CTG	TGG T 4- TCC AGG	ACT TGA T 40 TCT AGA	GAC-3' CTG-5' D TCC-3' AGG-5'
5'-GTG 3'-CAC V 5'-TGC 3'-ACG	CAG V 80	TCC AGG S Ppul 390 GGT	TAC ATG Y	CGG A	CAC V 400 GAC	GCT CGA A	CTC GAG L	AGC TCG S	TGT ACA C 10 ACC TGG	CAA GTT Q TGT ACA C	TGT ACA C 90	GCA CGT A 420 GAC	CTC GAG L	ACG C	GCG R 430 TTC	CGC GCG R	TCG S	TGG T 4. TCC	ACT TGA T 40 TCT AGA S	GAC-3' CTG-5' D TCC-3' AGG-5' S
5'-GTG 3'-CAC V 5'-TGC 3'-ACG	CAG V 80 GGG CCC	TCC AGG S Ppul 390 GGT CCA	TAC ATG Y	AAG TTC	CAC V 400 GAC CTG	GCT CGA A	CTC GAG L CCC GGG	AGC TCG S	TGT ACA C 10 ACC TGG	CAA GTT Q TGT ACA	TGT ACA C 90	GCA CGT A 420 GAC CTG	CTC GAG L CCC GGG	ACG C	GCG R 430 TTC AAG	CGC GCG R CAG GTC	TCG S GAC CTG	TGG T 4- TCC AGG	ACT TGA T 40 TCT AGA S	GAC-3' CTG-5' D TCC-3' AGG-5'
5'-GTG 3'-CAC V 5'-TGC 3'-ACG	CAG V 80 GGG CCC	TCC AGG S Ppul 390 GGT CCA	TAC ATG Y	AAG TTC	CAC V 400 GAC CTG	GCT CGA A	CTC GAG L CCC GGG	AGC TCG S	TGT ACA C 10 ACC TGG	CAA GTT Q TGT ACA C	TGT ACA C 90	GCA CGT A 420 GAC CTG	CTC GAG L CCC GGG	CGC GCG R	GCG R 430 TTC AAG . F	CGC GCG R CAG GTC	TCG S GAC CTG	TGG T 4- TCC AGG	ACT TGA T 40 TCT AGA S	GAC-3' CTG-5' D TCC-3' AGG-5' S
5'-GTG 3'-CAC V 5'-TGC 3'-ACG C	CAG V 80 GGG CCC G	TCC AGG S Ppul 390 GGT CCA	TAC ATG Y	AAG TTC	CAC V 400 GAC CTG	GCT CGA A	CTC GAG L CCC GGG	AGC TCG S 4 TTG AAC L	TGT ACA C 10 ACC TGG	TGT ACA C	TGT ACA C 90 GAT CTA D	GCA CGT A 420 GAC CTG	CTC GAG L CCC GGG	ACG C	GCG R 430 TTC AAG . F	CGC GCG R CAG GTC	GAC CTG D	TGG T 4- TCC AGG	ACT TGA T 40 TCT AGA S	GAC-3' CTG-5' D TCC-3' AGG-5' S
5'-GTG 3'-CAC V 5'-TGC 3'-ACG C	CAG V 80 GGG CCC G	Ppul 390 GGT CCA	TAC ATG Y	AAG TTC K	CAC V 400 GAC CTG D	GCT CGA A CAC GTG H	CCC GAG L CCC GGG P	AGC TCG S TTG AAC L	TGT ACA C	CAA GTT Q TGT ACA C	TGT ACA C 90 GAT CTA D	GCA CGT A 420 GAC CTG D	CCC GGG P	CGC GCG R	GCG R 430 TTC AAG F	CGC GCG R CAG GTC Q	GAC CTG D	TGG T TCC AGG S	ACT TGA T 40 TCT AGA S	GAC-3' CTG-5' D TCC-3' AGG-5' S
5'-GTG 3'-CAC V 5'-TGC 3'-ACG C 100	CAG V 80 	TCC AGG S Ppud 390 GGT CCA G	TAC ATG Y  GCC GGG P	AAG TTC K	CAC V 400 GAC CTG D	GCT CGA A CAC GTG H	CCC GGG P	AGC TCG S 4 TTG AAC L	3! TGT ACA C	CAA GTT Q TGT ACA C	TGT ACA C 90 GAT CTA D	GCA CGT A 420 GAC CTG D	CTC GAG L CCC GGG P	CGC GCG R Sma 490 CCG	GCG R 430 TTC AAG F AD	CGC GCG R CAG GTC Q GTC CCC	GAC CTG D	TGG T A TCC AGG S	ACT TGA T 40 TCT AGA S	GAC-3' CTG-5' D TCC-3' AGG-5' S 120
5'-GTG 3'-CAC V 5'-TGC 3'-ACG C 100	CAG V 80 GGG CCC G	TCC AGG S Ppud 390 GGT CCA G	TAC ATG Y  GCC GGG P	AAG TTC K	CAC V 400 GAC CTG D	GCT CGA A CAC GTG H	CCC GGG P	AGC TCG S 4 TTG AAC L	3! TGT ACA C	CAA GTT Q TGT ACA C	TGT ACA C 90 GAT CTA D	GCA CGT A 420 GAC CTG D	CTC GAG L CCC GGG P	CGC GCG R Sma 490 CCG	GCG R 430 TTC AAG F AD	CGC GCG R CAG GTC Q GTC CCC	GAC CTG D	TGG T A TCC AGG S	ACT TGA T 40 TCT AGA S	GAC-3' CTG-5' D TCC-3' AGG-5' S
5'-GTG 3'-CAC V 5'-TGC 3'-ACG C 100	CAG V 80 GGG CCC G	Ppul S Ppul 390 GGT CCA G	TAC ATG Y  41  CCC GGG P	AAG TTC K	CAC V 400 GAC CTG D	GCT CGA A CAC GTG H	CCC GGG P	AGC TCG S TTG AAC L	3! TGT ACA C	CAA GTT Q TGT ACA C 1110	TGT ACA C 90 GAT CTA D	GCA CGT A 420 GAC CTG D	CCC GGG P CTC GAG	ACG C CGC GCG R Sma 490 CCG GGC	430 TTC AAG F	CAG GTC Q CCC GGG	GAC CTG D	TGG T 4- TCC AGG S DO GAC CTC D	ACT TGA T T 40 TCT AGA S ACC TGG	GAC-3' CTG-5' D TCC-3' AGG-5' S 120
5'-GTG 3'-CAC V 5'-TGC 3'-ACG C 100	CAG V 80 GGG CCC G	Ppul S Ppul 390 GGT CCA G	TAC ATG Y  41  CCC GGG P	AAG TTC K	CAC V 400 GAC CTG D	GCT CGA A CAC GTG H	CCC GGG P	AGC TCG S TTG AAC L	35 TGT ACA C C 10 ACC TGG T	CAA GTT Q TGT ACA C 1110	TGT ACA C 90 GAT CTA D	GCA CGT A 420 GAC CTG D	CCC GGG P CTC GAG	CGC GCG R Sma 490 CCG GGC	430 TTC AAG F	CAG GTC Q CCC GGG	GAC CTG D	TGG T 4- TCC AGG S DO GAC CTC D	ACT TGA T  40 TCT AGA S  ACC TGG T	GAC-3' CTG-5' D TCC-3' AGG-5' S 120
5'-GTG 3'-CAC V 5'-TGC 3'-ACG C 100	CAG V 80 GGG CCC G	Ppul S Ppul 390 GGT CCA G	TAC ATG Y  41  CCC GGG P	AAG TTC K	CAC V 400 GAC CTG D	GCT CGA A CAC GTG H	CCC GGG P	AGC TCG S 4 TTG AAC L	35 TGT ACA C C 10 ACC TGG T	CAA GTT Q TGT ACA C 1110	TGT ACA C 90 GAT CTA D	GCA CGT A 420 GAC CTG D	CCC GGG P CTC GAG	CGC GCG R Sma 490 CCG GGC	430 TTC AAG F	CAG GTC Q CCC GGG	GAC CTG D	TGG T 4- TCC AGG S DO GAC CTC D	ACT TGA T  40 TCT AGA S  ACC TGG T	GAC-3' CTG-5' D TCC-3' AGG-5' S 120
5'-GTG 3'-CAC V 5'-TGC 3'-ACG C 100 5'-TCA 3'-AGT S	CAG V 80 GGG CCC G 450 AAG TTC	TCC AGG S Ppd 390 GGT CCA G GCC CGG A	TAC ATG Y  41  CCC GGG P  CCT GGA P	AAG TTC K	CAC V 400 GAC CTG D CCC GGG	GCT CGA A CAC GTG H	CCC GAG L CCC GGG P	AGC TCG S  4 TTG AAC L  70 CCA GGT P	35 TGT ACA C 10 ACC TGG T TCG S 130	CAA GTT Q TGT ACA C 110	TGT ACA C 90 GAT CTA D TCC AGG S	GCA CGT A 420 GAC CTG D	CCC GGG P CTC GAG	CGC GCG R Sma 490 CCG GGC	430 TTC AAG F	CAG GTC Q CCC GGG	GAC CTG D	TGG T 4- TCC AGG S DO GAC CTC D	ACT TGA T  40 TCT AGA S  ACC TGG T	GAC-3' CTG-5' D TCC-3' AGG-5' S 120
5'-GTG 3'-CAC V 5'-TGC 3'-ACG C 100 5'-TCA 3'-AGT S	CAG V 80 GGG CCC G AAG TTC K	TCC AGG S Ppul 390 GGT CCA GCC CGG A	TAC ATG Y  41  CCC GGG P  CCT GGA P	AAG TTC K	CCC GGG P	GCT CGA A CAC GTG H AGC TCG S	CTC GAG L CCC GGG P 4: CTT GAA L	AGC S 4 TTG AAC L CCA GGT P	TGT ACA C C TGG T T AGC S 130 539 CGC C	CAA GTT Q TGT ACA C 1110	TGT ACA C 90 GAT CTA D TCC AGG S	GCA CGT A 420 GAC CTG D	CCC GGG P CTC GAG	CGC GCG R Sma 490 CCG GGC	430 TTC AAG F	CAG GTC Q CCC GGG	GAC CTG D	TGG T 4- TCC AGG S DO GAC CTC D	ACT TGA T  40 TCT AGA S  ACC TGG T	GAC-3' CTG-5' D TCC-3' AGG-5' S 120
5'-TGC 3'-ACG C 100 5'-TCA 3'-AGT S	CAG V 80 GGG CCC G AAG TTC K	TCC AGG S Ppul 390 GGT CCA GCC CGG A	TAC ATG Y  41  CCC GGG P  CCT GGA P	AAG TTC K	CCC GGG P	GCT CGA A CAC GTG H AGC TCG S	CTC GAG L CCC GGG P 4: CTT GAA L	AGC S 4 TTG AAC L CCA GGT P	TGT ACA C C TGG T T AGC S 130 539 CGC C	CAA GTT Q TGT ACA C 110	TGT ACA C 90 GAT CTA D TCC AGG S	GCA CGT A 420 GAC CTG D	CCC GGG P CTC GAG	CGC GCG R Sma 490 CCG GGC	430 TTC AAG F	CAG GTC Q CCC GGG	GAC CTG D	TGG T 4- TCC AGG S DO GAC CTC D	ACT TGA T  40 TCT AGA S  ACC TGG T	GAC-3' CTG-5' D TCC-3' AGG-5' S 120

PCT/US91/03162 WO 91/16922

> 6/16 Figure 4b. Olionucleotides used to construct hCGB'

pKBM-hCGB - pKBM-hCGB'

Psti v v OxaNi(Bsu36i)

5'-G GGG GTG CTC CCG GCC CTG CC-3'

3'-AC GTC CCC CAC GAG GGC CGG GAC GGA GT-5'

LQGVLPALPQ

### 7/16 Figure 5. Oligonucleotides used to construct analogs F8, B9 and B11

5a Analog F8

PstI

OxaNI(Bsu361)

5'-G CCC CTG CC-3'

3'-AC GTC GGG GAC GGA GT-5'

L Q P L P. Q

45 46//51 52 53 54

5b Analog B9

ApaLI

Scal

1 Muq

5'-T GCA CTC TGC GAC AGC GAC AGT ACT GAC TGC GGG G-3'

3'-GAG ACG CTG TCG CTG TCA TGA CTG ACG CCC CCA G-5'

CALCDSDS TDCGGP

90

74 73 70 71

100

5c Analog B11;

Scal

StuI

Cassette A, A+ 5'-ACT GAC TGC ACC GTG AGA GGC CTC GGA CC-3'

A- 3'-TGA CTG ACG TGG CAC TCT CCG GAG CCT GGG AGA-5'

DCTVRGLGPS

100 ^ ^ ^ ^ ^ ^

Psti

BamHi

Cassette B, B+ 5'-C TCT TAC TGC AGC TTT GGT GAA ATG AAA GAA TAA G-3'

B- 3'-ATG ACG TCG AAA CCA CTT TAC TTT CTT ATT CCTAGG-5'

Y C S F G E M K E WWW

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Figure 6. Synthetic bovine alpha-subunit DNA and protein sequence

Eco	RI /	XhoI	_													
1		10			20			30				40			50	
5'-GA	TTCC	TCGA	GGGG	ATG	GAT	TAC	TAC	AGA	**	TAT	GCT	GCC	ATC	TTT	CTG	GTG-3'
3'-CT1	AAGG	AGCT	cccc	TAC												CAC-5'
				M	D	Y	Y	R	K	Y	A	A	I	F	L	V
				-24				-20								•
											0					
Dr	ilia <sup>.</sup> 06				70			80			90	MII.	-	11	00	
5/-404		AGT	GTG			CAT	GTT		CAT	TCC		CCG	GAT	-		TTT-3'
																AAA-5'
J .C.	L	s	٧	F	L	Н	V	L	Н	s	F	P	Đ	G	Ε	F
	-10									-1	1		•			
						8	sm!_									Fspll
	110			120				30			140			150		
																TCG-3'
			CCG		GGA P	CII	ACG C	TTC K	GAT L	K	E	IIG. N	K	ATG	AAA F	AGC-5'
T	M	Q	10	С	Ρ	=	·	•	_	•	-	П	20	•	r	3
_							_!	tsil						Xba	1	
_1	60			170			180			19	90		2	200		
		_														GCA-3'
•		-														CGT-5'
K	P	D	A	P	I	Y	Q	C	M	G	С	С	F	S	R	A (0
						30										40
					_Bg	111										
210			22	20	Bg		230			240			25	50		
5'-TAC	CCC		CCA	GCG	AGA	TCT	AAG			ATG			CCC	AAG		ATC-3'
5'-TAC 3'-ATG	CCC	TGA	CCA GGT	GCG	AGA TCT	TCT AGA	AAG TTC	TTC	TGT	ATG TAC	AAC	CAG	CCC	AAG TTC	TTG	TAG-5'
5'-TAC	CCC		CCA	GCG	AGA	TCT	AAG		TGT T	ATG			CCC	AAG		
5'-TAC 3'-ATG	CCC	TGA	CCA GGT	GCG CGC	AGA TCT	TCT AGA	AAG TTC	TTC	TGT	ATG TAC	AAC	CAG	CCC	AAG TTC	TTG	TAG-5'
5'-TAC 3'-ATG Y	CCC GGG P	TGA	CCA GGT	GCG CGC	AGA TCT	TCT AGA	AAG TTC K	TTC K	TGT T	ATG TAC	AAC	CAG	CCC	AAG TTC	TTG N	TAG-5'
5'-TAC 3'-ATG Y	CCC GGG P	TGA	CCA GGT	GCG CGC	AGA TCT	TCT AGA	AAG TTC K	TTC	TGT T 50	ATG TAC	AAC	CAG	CCC	AAG TTC	TTG	TAG-5'
5'-TAC 3'-ATG Y <u>Sp</u> 260	CCC GGG P	TGA T	CCA GGT P	GCG CGC A	AGA TCT R	TCT AGA S	AAG TTC K	TTC K Ball	TGT T 50	ATG TAC M	AAC L	CAG V	CCC GGG P	AAG TTC K	TTG N Bs1	TAG-5'
5'-TAC 3'-ATG Y <u>Sp</u> 260 5'-ACT	CCC GGG P	TGA T	CCA GGT P 270 GCT	GCG CGC A	AGA TCT R	TCT AGA S	AAG TTC K	TTC K Ball GCC	TGT T 50	ATG TAC M	AAC L	CAG V	CCC GGG P 300	AAG TTC K	TTG N BS1	TAG-5'
5'-TAC 3'-ATG Y <u>Sp</u> 260 5'-ACT	CCC GGG P	TGA T	CCA GGT P 270 GCT	GCG CGC A	AGA TCT R	TCT AGA S	AAG TTC K	TTC K Ball GCC	TGT T 50	ATG TAC M	AAC L	CAG V	CCC GGG P 300	AAG TTC K	TTG N BS1	TAG-5' I EXI GTG-3'
5'-TAC 3'-ATG Y <u>Sp</u> 260 5'-ACT 3'-TGA	CCC GGG P eI AGT TCA	TGA T	CCA GGT P 270 GCT CGA	GCG CGC A ACA TGT	AGA TCT R TGC ACG	TCT AGA S	AAG TTC K K SO GTG CAC	TTC K Ball GCC CGG	TGT T 50	ATG TAC M 290 GCA CGT	AAC L TTT AAA	CAG V ACC TGG	CCC GGG P 300 AAG TTC	AAG TTC K GCC CGG A	BS1 ACA TGT	TAG-5' I EXI GTG-3' CAC-5'
5'-TAC 3'-ATG Y <u>Sp</u> 260 5'-ACT 3'-TGA	CCC GGG P eI AGT TCA	TGA T GAA CTT E	CCA GGT P 270 GCT CGA	GCG CGC A ACA TGT	AGA TCT R TGC ACG	TCT AGA S	AAG TTC K K SO GTG CAC	TTC K Ball GCC CGG	TGT T 50 AAA TTT K	ATG TAC M	AAC L TTT AAA F	CAG V ACC TGG T	CCC GGG P 300 AAG TTC K	AAG TTC K GCC CGG A	N BS1	TAG-5' I EXI GTG-3' CAC-5'
5'-TAC 3'-ATG Y <u>Sp</u> 260 5'-ACT 3'-TGA	CCC GGG P eI AGT TCA	TGA T GAA CTT E	CCA GGT P 270 GCT CGA A	GCG CGC A ACA TGT	AGA TCT R TGC ACG	TCT AGA S 28 TGT ACA C	AAG TTC K SO GTG CAC V	TTC K Ball GCC CGG	TGT T 50 AAA TTT K	ATG TAC M 290 GCA CGT A	AAC L TTT AAA F	CAG V ACC TGG T	GCC GGG P 300 AAG TTC K	AAG TTC K GCC CGG A	BS1 ACA TGT	TAG-5' I EXI GTG-3' CAC-5' V
5'-TAC 3'-ATG Y 260 5'-ACT 3'-TGA T	GGG P eI AGT TCA S	TGA T GAA CTT E 60	CCA GGT P 270 GCT CGA A	GCG CGC A ACA TGT T	AGA TCT R TGC ACG C	TCT AGA S 28 TGT ACA C	AAG TTC K K SSO GTG CAC V	TTC K Ball GCC CGG A	TGT T 50	ATG TAC M 290 GCA CGT A	TTT AAA F	ACC TGG T	CCC GGG P 300 AAG TTC K	AAG TTC K GCC CGG A	BS1 ACA TGT T	TAG-5' I  EXI  GTG-3' CAC-5' V
5'-TAC 3'-ATG Y 260 5'-ACT 3'-TGA T	GGG P  AGT TCA S	GAA CTT E 60	CCA GGT P 270 GCT CGA A	GCG CGC A ACA TGT T	AGA TCT R TGC ACG C	TCT AGA S 28 TGT ACA C C 3330	AAG TTC K  BBO GTG CAC V	TTC K  Ball GCC CGG A	TGT T 50	ATG TAC M 290 GCA CGT A 26111	TTT AAA F	CAG V  ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst!	AAG TTC K GCC CGG A Sc	BSS ACA TGT T ACT	TAG-5' I  EXI  GTG-3' CAC-5' V  360 TGT-3'
5'-TAC 3'-ATG Y 260 5'-ACT 3'-TGA T	CCCC GGG P  eI  AGT TCA S  G GGG C CCT	TGA T GAA CTT E 60	CCA GGT P 270 GCT CGA A 320 GTC	GCG CGC A ACA TGT T	AGA TCT R TGC ACG C	TCT AGA S 28 TGT ACA C 3330 G GAG	AAG TTC K  E  GTG CAC V  AAG TTC TTC	TTC K  Ball GCC CGG A	TGT T 50 AAA TTT K	ATG TAC M 290 GCA CGT A 340 ; GAG	TTT AAA F	ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst! 350 : TGC	AAG TTC K GCC CGG A Sc AGT TCA	BS1 ACA TGT T AC1 TGGT T T AC1	TAG-5' I  CXI  GTG-3' CAC-5' V  - 360 TGT-3' ACA-5'
5'-TAC 3'-ATG Y 260 5'-ACT 3'-TGA T	CCCC GGG P  eI  AGT TCA S  G GGG C CCT	GAA CTT E 60	CCA GGT P 270 GCT CGA A	GCG CGC A ACA TGT T	AGA TCT R TGC ACG C	TCT AGA S 28 TGT ACA C C 3330	AAG TTC K  BBO GTG CAC V	TTC K  Ball GCC CGG A	TGT T 50	ATG TAC M 290 GCA CGT A 26111	TTT AAA F	CAG V  ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst!	AAG TTC K GCC CGG A Sc	BSS ACA TGT T ACT	TAG-5' I  EXI  GTG-3' CAC-5' V  360 TGT-3'
5'-TAC 3'-ATG Y 260 5'-ACT 3'-TGA T	CCCC GGG P  eI  AGT TCA S  G GGG C CCT	TGA T GAA CTT E 60	CCA GGT P 270 GCT CGA A 320 GTC	GCG CGC A ACA TGT T	AGA TCT R TGC ACG C	TCT AGA S 28 TGT ACA C 3330 G GAG	AAG TTC K  E  GTG CAC V  AAG TTC TTC	TTC K  Ball GCC CGG A	TGT T 50 AAA TTT K	ATG TAC M 290 GCA CGT A 340 ; GAG	TTT AAA F	ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst! 350 : TGC	AAG TTC K GCC CGG A Sc AGT TCA	BST ACA TGT T ACI TGG T	TAG-5' I  CXI  GTG-3' CAC-5' V  - 360 TGT-3' ACA-5'
5'-TAC 3'-ATG Y 260 5'-ACT 3'-TGA T	CCCC GGG P  eI  AGT TCA S  G GGG C CCT	TGA T GAA CTT E 60	CCA GGT P 270 GCT CGA A 320 GTC	GCG CGC A ACA TGT T	AGA TCT R TGC ACG C	TCT AGA S 28 TGT ACA C C 3330 C C C E	AAG TTC K  EB0 GTG CAC V	TTC K  Ball GCC CGG A	AAA TTT K  Or ACCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	ATG TAC M 290 GCA CGT A 340 GGGG GCA	TTT AAA F	ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst! 350 : TGC	AAG TTC K GCC CGG A Sc AGT TCA	BST ACA TGT T ACI TGG T	TAG-5' I  CXI  GTG-3' CAC-5' V  - 360 TGT-3' ACA-5'
5'-TAC 3'-ATG Y 260 5'-ACT 3'-TGA T	CCCC GGG P  eI  AGT TCA S  G GGG C CCT	TGA T GAA CTT E 60	CCA GGT P 270 GCT CGA A	GCG CGC A ACA TGT T	AGA TCT R TGC ACG C CAC	TCT AGA S 28 TGT ACA C C 3330 C C C E	AAG TTC K  EB0 GTG CAC V	TTC K  Ball GCC CGG A	TGT T SO AAA TTT K S ACC T T	ATG TAC M 290 GCA CGT A 340 GGGG GCA	TTT AAA F	ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst! 350 : TGC	AAG TTC K GCC CGG A Sc AGT TCA	BST ACA TGT T ACI TGG T	TAG-5' I  CXI  GTG-3' CAC-5' V  - 360 TGT-3' ACA-5'
5'-TAC 3'-ATG Y 260 5'-ACT 3'-TGA T 310 5'-ATI M	CCCC GGG P  eI  AGT TCA S  G GGG GGG CCT G	TGA T GAA CTT E 60	CCA GGT P 270 GCT CGA A 320 GTC CAG V	GCG CGC A ACA TGT T	AGA TCT R TGC ACG C CAC V 80	TCT AGA S 28 TGT ACA C 330 G GAG C CTC E SBEL 380 CCGC	AAG TTC K  E BO GTG CAC V  I AAG H I/BE	TTC K  GCC CGG A  CGC H  GCC GTC H  GTC H  GTC GTC GTC GTC GTC GTC GTC GTC GTC GT	TGT T 50  AAA TTT K  Dr 3  ACC ACC T T	ATG TAC M P P P P P P P P P P P P P P P P P P	TTT AAA F	ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst! 350 : TGC	AAG TTC K GCC CGG A Sc AGT TCA	BST ACA TGT T ACI TGG T	TAG-5' I  CXI  GTG-3' CAC-5' V  - 360 TGT-3' ACA-5'
5'-TAT 3'-ATG Y 260 5'-ACT 3'-TGA T 310 5'-ATA M	eI AGT TCA S G GGA TAT ATA	TGA T GAA CTT E 60 AAAT T TIA	CCA GGT P 270 GCT CGA A 320 CAG V	GCG CGC A ACA TGT T T TCT AGA	AGA TCT R TGC ACG C CAC V 80	TCT AGA S 28 TGT ACA C 330 G GAG C CTC E SBEL 380 CCGC	AAG TTC K  E BO GTG CAC V  I AAG H I/BE	TTC K  GCC CGG A  CGC H  GCC GTC H  GTC H  GTC GTC GTC GTC GTC GTC GTC GTC GTC GT	TGT T 50  AAA TTT K  Dr 3  ACC ACC T T	ATG TAC M P P P P P P P P P P P P P P P P P P	TTT AAA F	ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst! 350 : TGC	AAG TTC K GCC CGG A Sc AGT TCA	BST ACA TGT T ACI TGG T	TAG-5' I  CXI  GTG-3' CAC-5' V  - 360 TGT-3' ACA-5'
5'-TAC 3'-ATG Y 260 5'-ACT 3'-TGA T 310 5'-ATI M	CCCC GGG P  eI  AGT TCA S  G GGG GGG CCT G	TGA T GAA CTT E 60	CCA GGT P 270 GCT CGA A 320 GTC CAG V	GCG CGC A ACA TGT T T TCT AGA S	AGA TCT R TGC ACG C CAC V 80	TCT AGA S 28 TGT ACA C 330 G GAG C CTC E SBEL 380 CCGC	AAG TTC K  E BO GTG CAC V  I AAG H I/BE	TTC K  GCC CGG A  CGC H  GCC GTC H  GTC H  GTC GTC GTC GTC GTC GTC GTC GTC GTC GT	TGT T 50  AAA TTT K  Dr 3  ACC ACC T T	ATG TAC M P P P P P P P P P P P P P P P P P P	TTT AAA F	ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst! 350 : TGC	AAG TTC K GCC CGG A Sc AGT TCA	BST ACA TGT T ACI TGG T	TAG-5' I  CXI  GTG-3' CAC-5' V  - 360 TGT-3' ACA-5'
5'-TAT 3'-ATG Y 260 5'-ACT 3'-TGA T 310 5'-ATA M	eI AGT TCA S G GGA TAT ATA	TGA T GAA CTT E 60 AAAT T TIA	CCA GGT P 270 GCT CGA A 320 CAG V	GCG CGC A ACA TGT T T TCT AGA	AGA TCT R TGC ACG C CAC V 80	TCT AGA S 28 TGT ACA C 330 G GAG C CTC E SBEL 380 CCGC	AAG TTC K  E BO GTG CAC V  I AAG H I/BE	TTC K  GCC CGG A  CGC H  GCC GTC H  GTC H  GTC GTC GTC GTC GTC GTC GTC GTC GTC GT	TGT T 50  AAA TTT K  Dr 3  ACC ACC T T	ATG TAC M P P P P P P P P P P P P P P P P P P	TTT AAA F	ACC TGG T 70	CCC GGG P 300 AAG TTC K Pst! 350 : TGC	AAG TTC K GCC CGG A Sc AGT TCA	BST ACA TGT T ACI TGG T	TAG-5' I  CXI  GTG-3' CAC-5' V  - 360 TGT-3' ACA-5'

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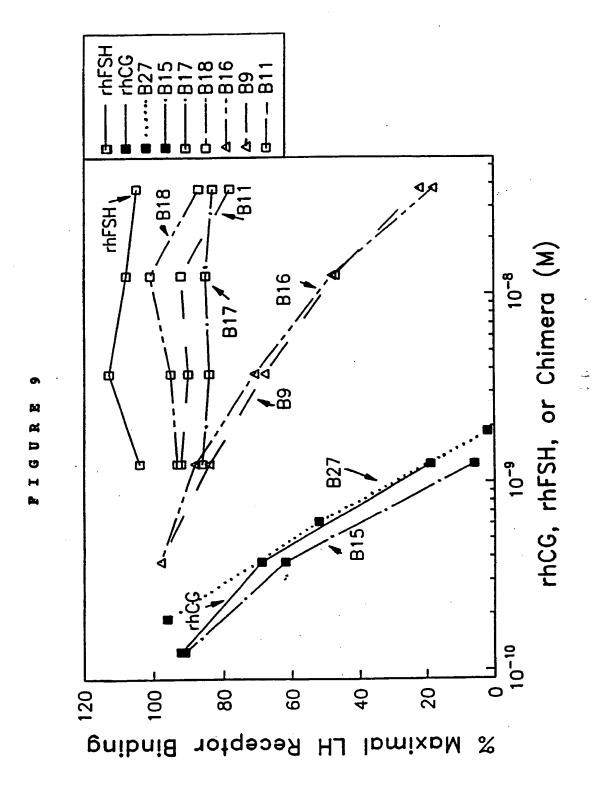
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Figure 7.	Regions of	human alpha-subunit	involved in	antibody binding

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	A101	,4109,45	01						
	/		/			A20	2,A112		
	A102	<b>!</b>			•	A10	9,A501		
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humen :	APDVQDCPECTLQEN	PFFSQPGAF	LACHGCCFS	KTMLVQKNVT	SESTCCVAKS	YNRVTVMGGF	KVENHTACHO	STCYYHKS	

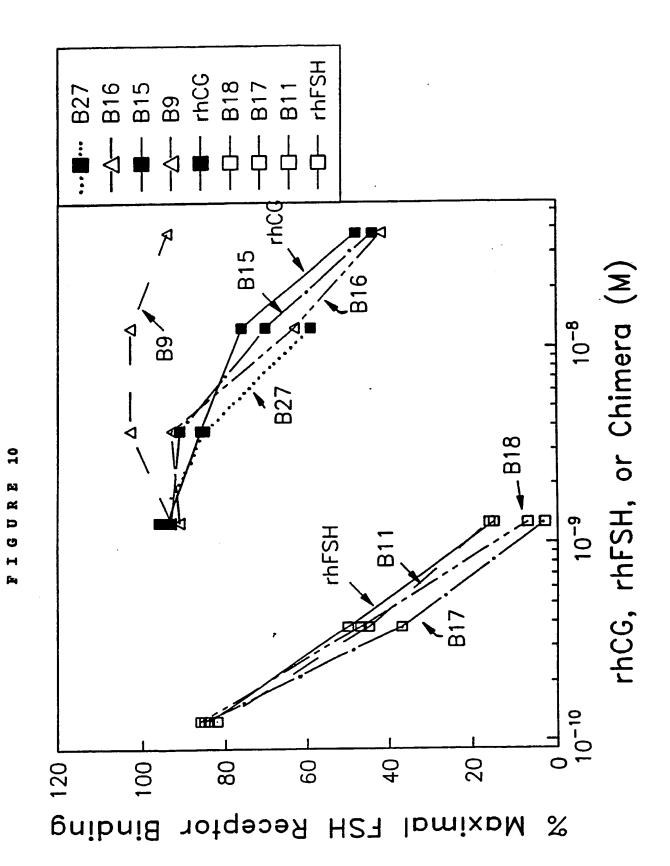
#### 10/16 FIGURE 8

hCG AND bLH RESIDUES WHICH DETERMINE ANTIBODY BINDING SPECIFICITY

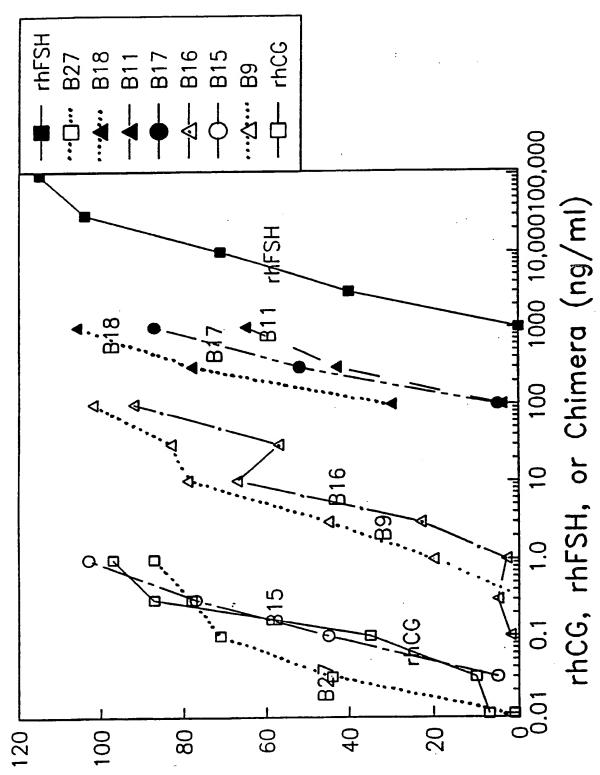
	B204 20
hCG hLH	SerLysGluProLeuArgProArgCysArgProIleAsnAlaThrLeuAlaValGluLys - Arg Trp - His Ile
	40
hCG hLH	GluGlyCysProValCysIleThrValAsnThrThrIleCysA1aGlyTyrCysProThr
FCG .	B101/B107/B109 60  MetThrArgValLeuGlnGlyValLeuProAlaLeuProGlnValValCysAsnTyrArg - Met Ala Pro Thr
	B108/B112 80
hCG hLH	AspValArgPheGluSerIleArgLeuProGlyCysProArgGlyValAsnProValVal
	B107/B109 · 100
hCG hLH	SerTyrAlaValAlaLeuSerCysGlnCysAlaLeuCysArgArgSerThrThrAspCys - PhePro Arg - GlyPro Ser
	B111 120
hCG hLH	GlyGlyProLysAspHisProLeuThrCysAspAspProArgPheGlnAspSerSerSerSer
	SCTP 140
hCG hLH	SerLysAlaProProProSerLeuProSerProSerArgLeuProGlyProSerAspThr Leu
hCG hLH	ProlieLeuProGin



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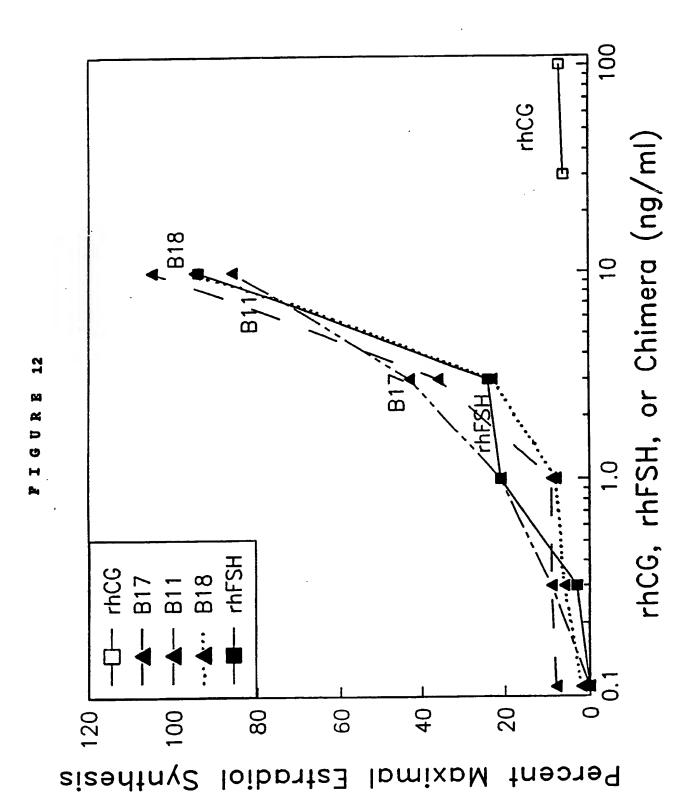


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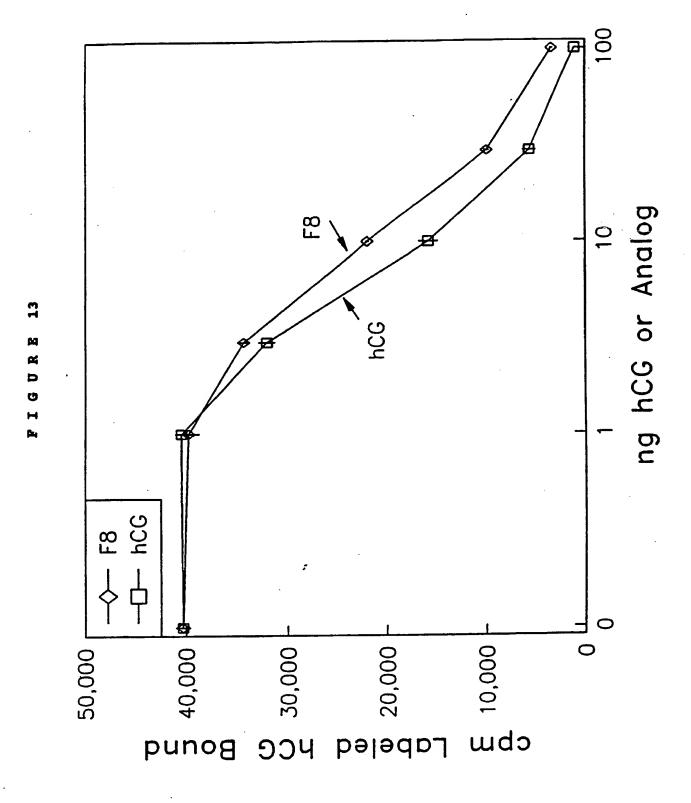


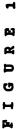
Percent Maximal Testosterone Synthesis

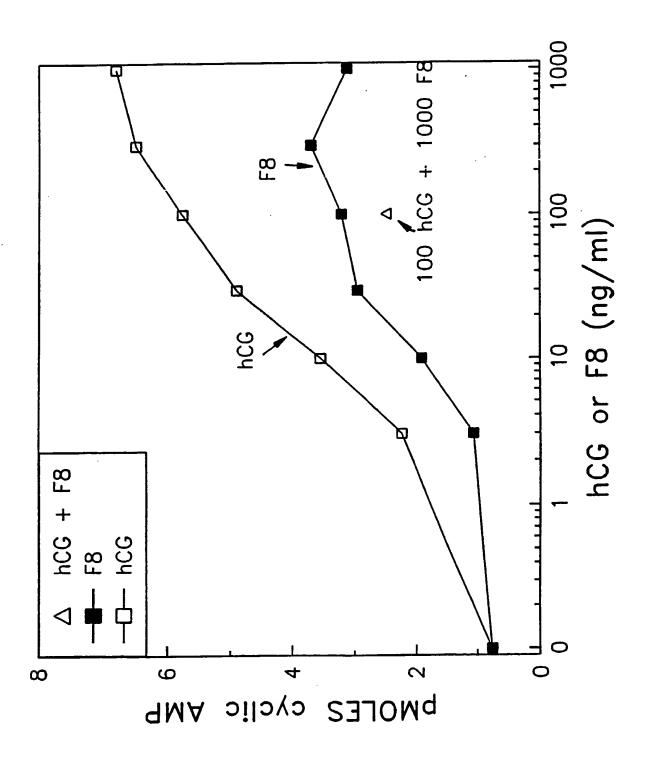
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#### INTERNATIONAL SEARCH REPORT

CLASS	SIFICATION OF SUBJECT MATTER HE CHARLES	PCF/U	<del>591/03162</del>
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	Cherionic Gonadotropin an	id Lutropin for	
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